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(54) Title: NOVEL THERAPEUTIC MOLECULAR VARIANTS AND USES THEREOF

(57) Abstract: The present invention relates generally to a sphingosine kinase variant and to derivatives, analogues, chemical equivalents and mimetics thereof exhibiting reduced catalytic activity and, more particularly, to sphingosine kinase variants which exhibit a reduced capacity to phosphorylate sphingosine to sphingosine-1-phosphate. The present invention also contemplates genetic sequences encoding said sphingosine kinase variants and derivatives, analogues and mimetics thereof. The variants of the present invention are useful in a range of therapeutic and prophylactic applications.

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NOVEL THERAPEUTIC MOLECULAR VARIANTS AND USES THEREOF

FIELD OF THE INVENTION

5 The present invention relates generally to a sphingosine kinase variant and to derivatives, analogues, chemical equivalents and mimetics thereof exhibiting reduced catalytic activity and, more particularly, to sphingosine kinase variants which exhibit a reduced capacity to phosphorylate sphingosine to sphingosine-1-phosphate. The present invention also
10 contemplates genetic sequences encoding said sphingosine kinase variants and derivatives, analogues and mimetics thereof. The variants of the present invention are useful in a range of therapeutic and prophylactic applications.

BACKGROUND OF THE INVENTION

15 Bibliographic details of the publications referred to by author in this specification are collected at the end of the description.

The reference to any prior art in this specification is not, and should not be taken as, an acknowledgment or any form of suggestion that prior art forms part of the common general
20 knowledge in Australia.

Sphingosine kinase is a key regulatory enzyme in a variety of cellular responses. Sphingosine-1-phosphate is known to be an important second messenger in signal transduction (Meyer *et al.*, 1997). It is mitogenic in various cell types (Alessenko, 1998)
25 and appears to trigger a diverse range of important regulatory pathways including prevention of ceramide-induced apoptosis (Culliver *et al.*, 1996), mobilisation of intracellular calcium by an IP₃-independent pathway, stimulation of DNA synthesis, activation of mitogen-activated protein (MAP) kinase pathway, activation of phospholipase D, and regulation of cell motility (for reviews see (Meyer *et al.*, 1997;
30 Spiegel *et al.*, 1998; Igarashi, 1997)).

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Recent studies (Xia *et al.*, 1998) have shown that sphingosine-1-phosphate is an obligatory signalling intermediate in the inflammatory response of vascular endothelial cells to tumour necrosis factor- α (TNF α). In spite of its obvious importance, very little is known of the mechanisms that control cellular sphingosine-1-phosphate levels. It is known that

5 sphingosine-1-phosphate levels in the cell are mediated largely by its formation from sphingosine by sphingosine kinase, and to a lesser extent by its degradation by endoplasmic reticulum-associated sphingosine-1-phosphate lyase and sphingosine-1-phosphate phosphatase (Spiegel *et al.*, 1998). Basal levels of sphingosine-1-phosphate in the cell are generally low, but can increase rapidly and transiently when cells are exposed

10 to mitogenic agents. This response appears correlated with an increase in sphingosine kinase activity in the cytosol and can be prevented by addition of the sphingosine kinase inhibitory molecules *N,N*-dimethylsphingosine and *DL-threo*-dihydrosphingosine. This indicates that sphingosine kinase is an important molecule responsible for regulating cellular sphingosine-1-phosphate levels. This places sphingosine kinase in a central and

15 obligatory role in mediating the effects attributed to sphingosine-1-phosphate in the cell.

Sphingosine kinase is speculated to play a role in a number of cellular activities including inflammation, calcium mobilisation, cell motility and adhesion molecule expression. Accordingly, there is a need to develop mechanisms of regulating these cellular activities

20 via regulation of the sphingosine kinase signalling pathway.

In work leading up to the present invention, the inventors have determined that amino acid sequence mutations introduced into the amino acid region defined by amino acid 16-153 of the human sphingosine kinase protein result in the production of a sphingosine kinase

25 variant which, in addition to exhibiting no sphingosine kinase baseline functional activity, also suppresses activation of wild-type sphingosine kinase molecules. Accordingly, the sphingosine kinase variants of the present invention both provide novel molecules for use in modulating sphingosine kinase signalling pathway function and facilitate the screening for and/or rational analysis, design and/or modification of agents for use in either

30 effectively mutating wild-type sphingosine kinase molecules or mimicing the activity of sphingosine kinase variant molecules.

SUMMARY OF THE INVENTION

Throughout this specification and the claims which follow, unless the context requires otherwise, the word "comprise", and variations such as "comprises" and "comprising", will
5 be understood to imply the inclusion of a stated integer or step or group of integers or steps but not the exclusion of any other integer or step or group of integers or steps.

Specific mutations in amino acid sequence are represented herein as "Xaa₁nXaa₂" where Xaa₁ is the original amino acid residue before mutation, n is the residue number and Xaa₂
10 is the mutant amino acid. The abbreviation "Xaa" may be the three letter or single letter amino acid code. A mutation in single letter code is represented, for example, by X₁nX₂ where X₁ and X₂ are the same as Xaa₁ and Xaa₂, respectively. The amino acid residues for sphingosine kinase are numbered with the residue glycine in the motif Asp Gly Leu Met (DGLM) being residue number 82.

15 The subject specification contains nucleotide and amino acid sequence information prepared using the programme PatentIn Version 2.0, presented herein after the bibliography. Each nucleotide or amino acid sequence is identified in the sequence listing by the numeric indicator <210> followed by the sequence identifier (e.g. <210>1, <210>2,
20 etc). The length, type of sequence (DNA, protein (PRT), etc) and source organism for each nucleotide or amino acid sequence are indicated by information provided in the numeric indicator fields <211>, <212> and <213>, respectively. Nucleotide and amino acid sequences referred to in the specification are defined by the information provided in numeric indicator field <400> followed by the sequence identifier (e.g. <400>1, <400>2,
25 etc).

One aspect of the present invention is directed to a sphingosine kinase variant comprising a mutation in a region defined by amino acids 16-153 or functionally equivalent region wherein said variant exhibits ablated or reduced catalytic activity relative to wild-type
30 sphingosine kinase or a derivative, homologue, analogue, chemical equivalent or mimetic of said sphingosine kinase variant.

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Another aspect of the present invention provides a human sphingosine kinase variant comprising a mutation in a region defined by amino acids 16-153 or functionally equivalent region wherein said variant exhibits ablated or reduced catalytic activity relative to wild-type human sphingosine kinase or a derivative, homologue, analogue, chemical
5 equivalent or mimetic of said sphingosine kinase variant.

In a preferred embodiment there is provided a human sphingosine kinase variant comprising an amino acid sequence with a single or multiple amino acid substitution, addition and/or deletion in a region defined by amino acids 16-153 or functionally
10 equivalent region wherein said variant exhibits ablated or reduced catalytic activity relative to wild-type sphingosine kinase or a derivative, homologue, analogue, chemical equivalent or mimetic of said sphingosine kinase variant.

In still a more preferred embodiment, there is provided a human sphingosine kinase variant comprising an amino acid sequence of the single or multiple amino acid substitution, addition and/or deletion in a region defined by amino acids 70-90, and more preferably 79-84, or functionally equivalent region wherein said variant exhibits ablated or reduced catalytic activity relative to wild-type sphingosine kinase or a derivative, homologue, analogue, chemical equivalent or mimetic of said sphingosine kinase variant.
20

In another preferred embodiment there is provided a human sphingosine kinase variant comprising an amino acid sequence with a single or multiple amino acid substitution, addition and/or deletion in a region defined by amino acid 16-153 or functionally equivalent region wherein said variant exhibits ablated catalytic activity relative to wild-
25 type sphingosine kinase or a derivative, homologue, analogue, chemical equivalent or mimetic of said sphingosine kinase variant.

In a most preferred embodiment, the subject sphingosine kinase variant comprises one or more of the amino acid substitutions selected from the following list:

- 5 -

- (i) G82D
- (ii) G82A
- (iii) G26D
- (iv) S79D
- 5 (v) G80D
- (vi) K103A
- (vii) G111D
- (viii) G113D
- (ix) G26A
- 10 (x) K27A
- (xi) K29A
- (xii) S79A
- (xiii) G80A
- (xiv) K103R
- 15 (xv) G111A

In another aspect the present invention is directed to a sphingosine kinase variant comprising a mutation in an ATP binding site region or functionally equivalent region wherein said variant exhibits ablated or reduced catalytic activity relative to wild-type sphingosine kinase or a derivative, homologue, analogue, chemical equivalent or mimetic of said sphingosine kinase variant .

Another aspect of the present invention is directed to an isolated nucleic acid molecule selected from the list consisting of:

- (i) An isolated nucleic acid molecule or derivative or equivalent thereof comprising a nucleotide sequence encoding or complementary to a sequence encoding a sphingosine kinase variant or derivative, homologue, analogue, chemical equivalent or mimetic of said variant which variant comprises a mutation in a region defined by amino acid 16-153 or functionally equivalent region wherein said variant

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exhibits ablated or reduced catalytic activity relative to wild-type sphingosine kinase.

- 5 (ii) An isolated nucleic acid molecule or derivative or equivalent thereof comprising a nucleotide sequence encoding or complementary to a sequence encoding a human sphingosine kinase variant or derivative, homologue, analogue, chemical equivalent or mimetic of said variant which variant comprises a mutation in a region defined by amino acid 16-153 or functionally equivalent region wherein said variant exhibits ablated or reduced catalytic activity relative to wild-type human
- 10 sphingosine kinase.
- (iii) An isolated nucleic acid molecule or derivative or equivalent thereof comprising a nucleotide sequence encoding or complementary to a sequence encoding a human sphingosine kinase variant or derivative, homologue, analogue, chemical equivalent
- 15 or mimetic of said variant, which variant comprises an amino acid sequence with a single or multiple multiple amino acid substitution, addition and/or deletion in a region defined by amino acid 16-153 or functionally equivalent region wherein said variant exhibits ablated or reduced catalytic activity relative to wild-type sphingosine kinase.
- 20 (iv) An isolated nucleic acid molecule or derivative or equivalent thereof comprising a nucleotide sequence encoding or complementary to a sequence encoding a human sphingosine kinase variant or derivative, homologue, analogue, chemical equivalent or mimetic of said variant, which variant comprises an amino acid sequence with a
- 25 single or multiple amino acid substitution, addition and/or deletion in a region defined by amino acid 70-90 or functionally equivalent region wherein said variant exhibits ablated or reduced catalytic activity relative to wild-type sphingosine kinase.
- 30 (v) An isolated nucleic acid molecule or derivative or equivalent thereof comprising a nucleotide sequence encoding or complementary to a sequence encoding a human

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5 sphingosine kinase variant or derivative, homologue, analogue, chemical equivalent or mimetic of said variant, which variant comprises an amino acid sequence with a single or multiple multiple amino acid substitution, addition and/or deletion in a region defined by amino acid 79-84 or functionally equivalent region wherein said variant exhibits ablated or reduced catalytic activity relative to wild-type sphingosine kinase.

10 (vi) An isolated nucleic acid molecule or derivative or equivalent thereof comprising a nucleotide sequence encoding or complementary to a sequence encoding a sphingosine kinase variant or a derivative, homologue, analogue, chemical equivalent or mimetic of said variant comprising one or more of the amino acid substitutions selected from the following list:

- 15 (a) G82D
(b) G82A
(c) G26D
(d) S79D
(e) G80D
(f) K103A
(g) G111D
20 (h) G113D
(i) G26A
(j) K27A
(k) K29A
(l) S79A
25 (m) G80A
(n) K103R
(o) G111A

30 (vii) An isolated nucleic acid molecule or derivative or analogue thereof comprising a nucleotide sequence encoding or complementary to a sequence encoding a sphingosine kinase variant or derivative, homologue, analogue, chemical equivalent

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or mimetic of said variant which variant comprises a mutation in an ATP binding site region or functionally equivalent region wherein said variant exhibits ablated or reduced catalytic activity relative to wild-type sphingosine kinase.

5 Accordingly, another aspect of the present invention provides a method for detecting an agent capable of modulating the interaction of FOSK with sphingosine kinase or its functional equivalent or derivative thereof said method comprising contacting a cell or extract thereof containing said sphingosine kinase and FOSK or its functional equivalent or derivative with a putative agent and detecting an altered expression phenotype associated
10 with said interaction.

In yet another aspect the present invention provides a method for detecting an agent capable of binding or otherwise associating with the sphingosine kinase region defined by amino acids 16-153 or functional equivalent or derivative thereof said method comprising
15 contacting a cell containing said amino acid region or functional equivalent or derivative thereof with a putative agent and detecting an altered expression phenotype associated with modulation of the function of sphingosine kinase or its functional equivalent or derivative.

Accordingly, another aspect of the present invention is directed to a method for analysing,
20 designing and/or modifying an agent capable of interacting with the sphingosine kinase region defined by amino acids 16-153 or derivative thereof and modulating at least one functional activity associated with said sphingosine kinase said method comprising contacting said sphingosine kinase or derivative thereof with a putative agent and assessing the degree of interactive complementarity of said agent with said binding site.

25

In a related aspect, the present invention should be understood to extend to the agents identified utilising any of the methods hereinbefore defined. In this regard, reference to an agent should be understood as a reference to any proteinaceous or non-proteinaceous molecule which modulates at least one sphingosine kinase functional activity.

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Another aspect of the present invention contemplates a method of modulating cellular functional activity in a mammal said method comprising administering to said mammal an effective amount of a sphingosine kinase variant or agent as hereinbefore defined for a time and under conditions sufficient to inhibit, reduce or otherwise down-regulate at least
5 one functional activity of wild-type sphingosine kinase.

Another aspect of the present invention relates to the treatment and/or prophylaxis of a condition in a mammal, which condition is characterised by aberrant, unwanted or otherwise inappropriate cellular activity, said method comprising administering to said
10 mammal an effective amount of a sphingosine kinase variant or agent as hereinbefore defined for a time and under conditions sufficient to inhibit, reduce or otherwise down-regulate at least one functional activity of wild-type sphingosine kinase wherein said down-regulation results in modulation of cellular functional activity.

15 A further aspect of the present invention relates to the use of a sphingosine kinase variant or agent as hereinbefore defined in the manufacture of a medicament for the modulation of cellular functional activity.

Another aspect of the present invention relates to a sphingosine kinase variant or agent as
20 hereinbefore defined for use in modulating cellular functional activity.

In yet another further aspect the present invention contemplates a pharmaceutical composition comprising a sphingosine kinase variant or agent as hereinbefore defined together with one or more pharmaceutically acceptable carriers and/or diluents.
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Single and three letter abbreviations used throughout the specification are defined in Table 1.

TABLE 1

5 **Single and three letter amino acid abbreviations**

	Amino Acid	Three-letter Abbreviation	One-letter Symbol
10	Alanine	Ala	A
	Arginine	Arg	R
	Asparagine	Asn	N
	Aspartic acid	Asp	D
	Cysteine	Cys	C
15	Glutamine	Gln	Q
	Glutamic acid	Glu	E
	Glycine	Gly	G
	Histidine	His	H
	Isoleucine	Ile	I
20	Leucine	Leu	L
	Lysine	Lys	K
	Methionine	Met	M
	Phenylalanine	Phe	F
	Proline	Pro	P
25	Serine	Ser	S
	Threonine	Thr	T
	Tryptophan	Trp	W
	Tyrosine	Tyr	Y
	Valine	Val	V
30	Any residue	Xaa	X

BRIEF DESCRIPTION OF THE DRAWINGS

Figure 1 is a schematic representation of the sequence alignment of the putative catalytic domains of some diacylglycerol kinases with sphingosine kinases. Highly conserved residues within the putative catalytic domain of diacylglycerol kinases are highlighted.
5 The marked (•) residue indicates the site where mutagenesis (Gly → Asp) in these three diacylglycerol kinases ablates catalytically activity.

Figure 2 is both a graphical representation and image of site directed mutagenesis of
10 human sphingosine kinase HEK293 cells transfected with either pcDNA3-SK, pcDNA3-G26DSK, pcDNA3-S79DSK, pcDNA3-G80DSK, pcDNA3-G82DSK, pcDNA3-K103ASK, pcDNA3-G111DSK, pcDNA3-G113DSK, or empty pcDNA3 vector were harvested and analysed for (A) protein expression levels by Western blot using the M2 anti-FLAG antibody, and (B) sphingosine kinase activity.

15 **Figure 3** is a graphical representation demonstrating that expression of G82D SK in HEK293 cells blocks activation of endogenous sphingosine kinase activity by TNF α , PMA and IL-1. HEK293 cells transfected with either pcDNA3-G82DSK or empty pcDNA3 vector were treated with 1ng/ml TNF α and 100 units/ml IL-1 for 10 min and 100 ng/ml
20 PMA for 30 min.

Figure 4 is a graphical representation of time course of sphingosine kinase activation by TNF α in HEK293 cells expressing G82D SK. HEK293 cells transfected with either pcDNA3-G82DSK or empty pcDNA3 vector were treated with 1ng/ml TNF α . Cells were
25 harvested at various times over 45 min. of TNF α treatment with the cell lysates assayed for sphingosine kinase activity.

Figure 5 is a graphical representation demonstrating that expression of G82D SK in HEK293 cells decreases activation of overexpressed wild-type sphingosine kinase activity
30 by TNF α . HEK293 cells, either transfected with pcDNA3-SK, or cotransfected with equal proportions of pcDNA3-SK and pcDNA3-G82DSK were treated with 1 ng/ml

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TNF α for 10 min. Cells were then harvested and the SK activity in the cell lysates determined.

Figure 6 is a graphical representation that the expression of G82D SK in 3T3 fibroblasts
5 inhibiting activation of SK by the oncogene Ras.

Figure 7 is a graphical representation that the expression of G82D SK in HEK293 cells does not effect activation of protein kinase C activity by TNF α or PMA.

10 **Figure 8** is an image demonstrating that the expression of G82D SK in HEK293 cells does not inhibit activation of sphingomyelinase by TNF.

Figure 9 is a graphical representation demonstrating that the expression of G82D SK in HEK293 prevents ERK activation by TNF α .

15

Figure 10 is a schematic representation of the types of drugs which can be identified and/or developed in light of the development of the present invention.

Figure 11 is a graphical representation demonstrating that G82D SK inhibits Ras
20 transformation. A. NIH 3T3 cells were transfected with V12-Ras, v-Src or V12-Ras plus G82D-SK, SphK activity was measured 48 h after transfection. B. Focus formation assays were performed in V12-Ras, v-Src, SphK, or V12-Ras plus G82D-SK transfected NIH 3T3 cells in the absence or presence of DMS (2.5 μ M) over two weeks.

25 **Figure 12** shows site directed mutagenesis of human sphingosine kinase HEK293T cells transfected with either pcDNA3-SK^{WT}, pcDNA3-SK^{G82D}, pcDNA3-SK^{G82A}, or empty pcDNA3 vector were harvested and analysed for (A) protein expression levels by Western blot using the MT anti-FLAG antibody, and (B) sphingosine kinase activity.

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Figure 13 shows kinetic analysis with ATP of (A) hSK^{WT} and (B) hSK^{G82A}. Kinetic analyses were performed with ATP in the concentration range of 0-2 mM and 0-40 mM for hSK^{WT} and hSK^{G82A}, respectively. In both cases sphingosine was present at 100 μ M.

- 5 **Figure 14** shows kinetic analysis with sphingosine of (A) hSK^{WT} and (B) hSK^{G82A}. Kinetic analyses were performed with sphingosine in the concentration range of 0-2 mM for both hSK^{WT} and hSK^{G82A}. ATP was present at 1 mM and 40 mM for hSK^{WT} and hSK^{G82A}, respectively.
- 10 **Figure 15** is a graphical representation of site directed mutagenesis of human sphingosine kinase. HEK293T cells transfected with either empty pcDNA3 vector, pcDNA3-SK^{WT}, or pcDNA3-mutant hSK were harvested and analysed for sphingosine kinase activity.

DETAILED DESCRIPTION OF THE PREFERRED EMBODIMENTS

The present invention is predicated, in part, on the determination that the ablation of catalytic activity of human sphingosine kinase can be achieved by introducing a mutation
5 into the amino acid region defined by amino acids 16-153. Further, the introduction of such a mutation not only generates a sphingosine kinase variant which exhibits ablated, reduced or a limited baseline functional activity but also generates a variant which can function as a dominant negative sphingosine kinase, either *in vitro* or *in vivo*, in that it inhibits the activation of wild-type sphingosine kinase. This determination facilitates the
10 rational design of products and methodology for use in the therapy and prophylaxis of conditions characterised by the aberrant, unwanted or otherwise inappropriate functioning of sphingosine kinase signalling.

Accordingly, one aspect of the present invention is directed to a sphingosine kinase variant
15 comprising a mutation in a region defined by amino acids 16-153 or functionally equivalent region wherein said variant exhibits ablated or reduced catalytic activity relative to wild-type sphingosine kinase or a derivative, homologue, analogue, chemical equivalent or mimetic of said sphingosine kinase variant .

20 Reference to "sphingosine kinase" should be understood as including a reference to all forms of sphingosine kinase protein or derivatives, homologues, analogues, equivalents or mimetics thereof. In this regard, "sphingosine kinase" should be understood as being a molecule which is, *inter alia*, involved in the generation of sphingosine-1-phosphate during activation of the sphingosine kinase signalling pathway. This includes, for
25 example, all protein forms of sphingosine kinase or its functional derivatives, homologues, analogues, equivalents or mimetics thereof including, for example, any isoforms which arise from alternative splicing of sphingosine kinase mRNA or allelic or polymorphic variants of sphingosine kinase. Preferably, said sphingosine kinase is human sphingosine kinase.

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Accordingly there is more particularly provided a human sphingosine kinase variant comprising a mutation in a region defined by amino acids 16-153 or functionally equivalent region wherein said variant exhibits ablated or reduced catalytic activity relative to wild-type human sphingosine kinase or a derivative, homologue, analogue, chemical
5 equivalent or mimetic of said sphingosine kinase variant .

The term "protein" should be understood to encompass peptides, polypeptides and proteins. The protein may be glycosylated or unglycosylated and/or may contain a range of other molecules fused, linked, bound or otherwise associated to the protein such as
10 amino acids, lipids, carbohydrates or other peptides, polypeptides or proteins. Reference hereinafter to a "protein" includes a protein comprising a sequence of amino acids as well as a protein associated with other molecules such as amino acids, lipids, carbohydrates or other peptides, polypeptides or proteins.

15 Reference to "mutation" should be understood as a reference to any change, alteration or other modification, whether occurring naturally or non-naturally, which renders a sphingosine kinase molecule catalytically inactive or capable only of a reduced level of catalytic activity. In this regard, the phrase "catalytic activity" in the context of sphingosine kinase activity should be understood as a reference to the capacity of
20 sphingosine kinase to phosphorylate sphingosine to sphingosine-1 phosphate.

The change, alteration or other modification may take any form including, but not limited to, a structural modification (such an alteration in the secondary, tertiary or quaternary structure of the sphingosine kinase molecule), a molecular modification (such as an
25 addition, substitution or deletion of one or more amino acids from the sphingosine kinase protein) or a chemical modification. The subject modification should also be understood to extend to the fusion, linking or binding of a proteinaceous or non-proteinaceous molecule to the sphingosine kinase protein or to the nucleic acid molecule encoding a sphingosine kinase protein thereby rendering the expression product either catalytically
30 inactive or capable only of reduced catalytic activity. It should also be understood that although it is necessary that the subject mutation is expressed by the sphingosine kinase

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expression product, the creation of the mutation may be achieved by any suitable means including either mutating a wild-type sphingosine kinase protein, synthesising a sphingosine kinase variant or modifying a nucleic acid molecule encoding a wild-type sphingosine kinase protein such that the expression product of said mutated nucleic acid
5 molecule is a sphingosine kinase protein variant. Preferably, said mutation is a single or multiple amino acid sequence substitution, addition and/or deletion.

In accordance with this preferred embodiment there is provided a human sphingosine kinase variant comprising an amino acid sequence with a single or multiple amino acid
10 substitution, addition and/or deletion in a region defined by amino acids 16-153 or functionally equivalent region wherein said variant exhibits ablated or reduced catalytic activity relative to wild-type sphingosine kinase or a derivative, homologue, analogue, chemical equivalent or mimetic of said sphingosine kinase variant .

15 In terms of the present invention, reference to "wild-type" sphingosine kinase is a reference to the forms of sphingosine kinase expressed by most individuals in a given population wherein the subject sphingosine kinase is catalytically active within the context discussed hereinbefore. There may be greater than one wild-type form of sphingosine kinase (for example due to allelic or isoform variation) and the level of catalytic activity exhibited by
20 said wild-type sphingosine kinase molecules may fall within a range of levels. However, it should be understood that "wild-type" does not include reference to a naturally occurring form of sphingosine kinase which is not catalytically active. Such a variant form of sphingosine kinase may, in fact, constitute a naturally occurring mutant form of sphingosine kinase within the context of the present invention.

25

In still a more preferred embodiment, there is provided a human sphingosine kinase variant comprising an amino acid sequence of the single or multiple amino acid substitution, addition and/or deletion in a region defined by amino acids 70-90, and more preferably 79-84, or functionally equivalent region wherein said variant exhibits ablated or reduced
30 catalytic activity relative to wild-type sphingosine kinase or a derivative, homologue, analogue, chemical equivalent or mimetic of said sphingosine kinase variant.

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In a most preferred embodiment, the subject sphingosine kinase variant comprises an amino acid substitution of the glycine amino acid at position 82 to aspartic acid.

Without limiting the invention to any one theory or mode of action, sphingosine kinase is
5 thought to exhibit two levels of catalytic activity. At the first level, sphingosine kinase
exhibits baseline catalytic activity. At the second level, sphingosine kinase exhibiting
baseline activity can be activated such that the Vmax of the enzyme is increased. In the
context of the present invention, the ablation or reduction of sphingosine kinase catalytic
activity will be achieved where the baseline activity and/or the activation of sphingosine
10 kinase beyond that of baseline activity is ablated or reduced. Preferably, both levels of
activity are ablated or reduced and even more preferably both levels of activity are ablated.

In another preferred embodiment there is provided a human sphingosine kinase variant
comprising an amino acid sequence with a single or multiple amino acid substitution,
15 addition and/or deletion in a region defined by amino acid 16-153 or functionally
equivalent region wherein said variant exhibits ablated catalytic activity relative to wild-
type sphingosine kinase or a derivative, homologue, analogue, chemical equivalent or
mimetic of said sphingosine kinase variant.

20 Preferably said subject human sphingosine kinase variant comprises an amino acid
addition, substitution and/or deletion in the region defined by amino acids 70-90 and even
more preferably 79-84.

In a most preferred embodiment, the subject sphingosine kinase variant comprises one or
25 more of the amino acid substitutions selected from the following list:

- (i) G82D
- (ii) G82A
- (iii) G26D
- (iv) S79D
- 30 (v) G80D
- (vi) K103A

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- (vii) G111D
- (viii) G113D
- (ix) G26A
- (x) K27A
- 5 (xi) K29A
- (xii) S79A
- (xiii) G80A
- (xiv) K103R
- (xv) G111A

10

"Derivatives" include fragments, parts, portions, variants and mimetics from natural, synthetic or recombinant sources including fusion proteins. Parts or fragments include, for example, active regions of sphingosine kinase. Derivatives may be derived from insertion, deletion or substitution of amino acids. Amino acid insertional derivatives include amino and/or carboxylic terminal fusions as well as intrasequence insertions of single or multiple amino acids. Insertional amino acid sequence variants are those in which one or more amino acid residues are introduced into a predetermined site in the protein although random insertion is also possible with suitable screening of the resulting product.

15

Deletional variants are characterized by the removal of one or more amino acids from the sequence. Substitutional amino acid variants are those in which at least one residue in the sequence has been removed and a different residue inserted in its place. An example of substitutional amino acid variants are conservative amino acid substitutions. Conservative amino acid substitutions typically include substitutions within the following groups: glycine and alanine; valine, isoleucine and leucine; aspartic acid and glutamic acid; asparagine and glutamine; serine and threonine; lysine and arginine; and phenylalanine and tyrosine. Additions to amino acid sequences include fusions with other peptides, polypeptides or proteins.

25

Reference to "homologues" should be understood as a reference to sphingosine kinase nucleic acid molecules or proteins derived from species other than the species being treated.

30

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Chemical and functional equivalents of sphingosine kinase nucleic acid or protein molecules should be understood as molecules exhibiting any one or more of the functional activities of these molecules and may be derived from any source such as being chemically synthesized or identified via screening processes such as natural product screening.

5

The derivatives include fragments having particular epitopes or parts of the entire protein fused to peptides, polypeptides or other proteinaceous or non-proteinaceous molecules.

10 Analogues contemplated herein include, but are not limited to, modification to side chains, incorporating of unnatural amino acids and/or their derivatives during peptide, polypeptide or protein synthesis and the use of crosslinkers and other methods which impose conformational constraints on the proteinaceous molecules or their analogues.

15 Derivatives of nucleic acid sequences may similarly be derived from single or multiple nucleotide substitutions, deletions and/or additions including fusion with other nucleic acid molecules. The derivatives of the nucleic acid molecules of the present invention include oligonucleotides, PCR primers, antisense molecules, molecules suitable for use in cosuppression and fusion of nucleic acid molecules. Derivatives of nucleic acid sequences also include degenerate variants.

20

Examples of side chain modifications contemplated by the present invention include modifications of amino groups such as by reductive alkylation by reaction with an aldehyde followed by reduction with NaBH_4 ; amidination with methylacetimidate; acylation with acetic anhydride; carbamoylation of amino groups with cyanate; 25 trinitrobenzylation of amino groups with 2, 4, 6-trinitrobenzene sulphonic acid (TNBS); acylation of amino groups with succinic anhydride and tetrahydrophthalic anhydride; and pyridoxylation of lysine with pyridoxal-5-phosphate followed by reduction with NaBH_4 .

30 The guanidine group of arginine residues may be modified by the formation of heterocyclic condensation products with reagents such as 2,3-butanedione, phenylglyoxal and glyoxal.

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The carboxyl group may be modified by carbodiimide activation *via* O-acylisourea formation followed by subsequent derivitisation, for example, to a corresponding amide.

Sulphydryl groups may be modified by methods such as carboxymethylation with
5 iodoacetic acid or iodoacetamide; performic acid oxidation to cysteic acid; formation of a mixed disulphides with other thiol compounds; reaction with maleimide, maleic anhydride or other substituted maleimide; formation of mercurial derivatives using 4-chloromercuribenzoate, 4-chloromercuriphenylsulphonic acid, phenylmercury chloride, 2-chloromercuri-4-nitrophenol and other mercurials; carbamoylation with cyanate at alkaline
10 pH.

Tryptophan residues may be modified by, for example, oxidation with N-bromosuccinimide or alkylation of the indole ring with 2-hydroxy-5-nitrobenzyl bromide or sulphenyl halides. Tyrosine residues on the other hand, may be altered by nitration with
15 tetranitromethane to form a 3-nitrotyrosine derivative.

Modification of the imidazole ring of a histidine residue may be accomplished by alkylation with iodoacetic acid derivatives or N-carboethoxylation with diethylpyrocarbonate.

20 Examples of incorporating unnatural amino acids and derivatives during protein synthesis include, but are not limited to, use of norleucine, 4-amino butyric acid, 4-amino-3-hydroxy-5-phenylpentanoic acid, 6-aminohexanoic acid, t-butylglycine, norvaline, phenylglycine, ornithine, sarcosine, 4-amino-3-hydroxy-6-methylheptanoic acid, 2-thienyl
25 alanine and/or D-isomers of amino acids. A list of unnatural amino acids contemplated herein is shown in Table 2.

TABLE 2

	Non-conventional amino acid	Code	Non-conventional amino acid	Code
5	α -aminobutyric acid	Abu	L-N-methylalanine	Nmala
	α -amino- α -methylbutyrate	Mgab	L-N-methylarginine	Nmarg
	aminocyclopropane-	Cpro	L-N-methylasparagine	Nmasn
10	carboxylate		L-N-methylaspartic acid	Nmasp
	aminoisobutyric acid	Aib	L-N-methylcysteine	Nmcys
	aminonorbornyl-	Norb	L-N-methylglutamine	Nmgln
	carboxylate		L-N-methylglutamic acid	Nmglu
	cyclohexylalanine	Chexa	L-N-methylhistidine	Nmhis
15	cyclopentylalanine	Cpen	L-N-methylisoleucine	Nmile
	D-alanine	Dal	L-N-methylleucine	Nmleu
	D-arginine	Darg	L-N-methyllysine	Nmlys
	D-aspartic acid	Dasp	L-N-methylmethionine	Nmmet
	D-cysteine	Dcys	L-N-methylnorleucine	Nmnle
20	D-glutamine	Dgln	L-N-methylnorvaline	Nmnva
	D-glutamic acid	Dglu	L-N-methylornithine	Nmorn
	D-histidine	Dhis	L-N-methylphenylalanine	Nmphe
	D-isoleucine	Dile	L-N-methylproline	Nmpro
	D-leucine	Dleu	L-N-methylserine	Nmser
25	D-lysine	Dlys	L-N-methylthreonine	Nmthr
	D-methionine	Dmet	L-N-methyltryptophan	Nmtrp
	D-ornithine	Dorn	L-N-methyltyrosine	Nmtyr
	D-phenylalanine	Dphe	L-N-methylvaline	Nmval
	D-proline	Dpro	L-N-methylethylglycine	Nmetg
30	D-serine	Dser	L-N-methyl-t-butylglycine	Nmtbug
	D-threonine	Dthr	L-norleucine	Nle
	D-tryptophan	Dtrp	L-norvaline	Nva
	D-tyrosine	Dtyr	α -methyl-aminoisobutyrate	Maib

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	D-valine	Dval	α -methyl- -aminobutyrate	Mgab
	D- α -methylalanine	Dmala	α -methylcyclohexylalanine	Mchexa
	D- α -methylarginine	Dmarg	α -methylcyclopentylalanine	Mcpen
	D- α -methylasparagine	Dmasn	α -methyl- α -naphthylalanine	Manap
5	D- α -methylaspartate	Dmasp	α -methylpenicillamine	Mpen
	D- α -methylcysteine	Dmcys	N-(4-aminobutyl)glycine	Nglu
	D- α -methylglutamine	Dmgln	N-(2-aminoethyl)glycine	Naeg
	D- α -methylhistidine	Dmhis	N-(3-aminopropyl)glycine	Norn
	D- α -methylisoleucine	Dmile	N-amino- α -methylbutyrate	Nmaabu
10	D- α -methylleucine	Dmleu	α -naphthylalanine	Anap
	D- α -methyllysine	Dmlys	N-benzylglycine	Nphe
	D- α -methylmethionine	Dmmet	N-(2-carbamylethyl)glycine	Ngln
	D- α -methylornithine	Dmorn	N-(carbamylmethyl)glycine	Nasn
	D- α -methylphenylalanine	Dmphe	N-(2-carboxyethyl)glycine	Nglu
15	D- α -methylproline	Dmpro	N-(carboxymethyl)glycine	Nasp
	D- α -methylserine	Dmser	N-cyclobutylglycine	Ncbut
	D- α -methylthreonine	Dmthr	N-cycloheptylglycine	Nchep
	D- α -methyltryptophan	Dmtrp	N-cyclohexylglycine	Nchex
	D- α -methyltyrosine	Dmtty	N-cyclodecylglycine	Ndec
20	D- α -methylvaline	Dmval	N-cyclododecylglycine	Ncdod
	D-N-methylalanine	Dnmala	N-cyclooctylglycine	Ncoct
	D-N-methylarginine	Dnmarg	N-cyclopropylglycine	Ncpro
	D-N-methylasparagine	Dnmasn	N-cycloundecylglycine	Ncund
	D-N-methylaspartate	Dnmasp	N-(2,2-diphenylethyl)glycine	Nbhm
25	D-N-methylcysteine	Dnmcys	N-(3,3-diphenylpropyl)glycine	Nbhe
	D-N-methylglutamine	Dnmgln	N-(3-guanidinopropyl)glycine	Narg
	D-N-methylglutamate	Dnmglu	N-(1-hydroxyethyl)glycine	Nthr
	D-N-methylhistidine	Dnmhis	N-(hydroxyethyl)glycine	Nser
	D-N-methylisoleucine	Dnmile	N-(imidazolylethyl)glycine	Nhis
30	D-N-methylleucine	Dnmleu	N-(3-indolylyethyl)glycine	Nhtrp
	D-N-methyllysine	Dnmlys	N-methyl- γ -aminobutyrate	Nmgabu
	N-methylcyclohexylalanine	Nmchexa	D-N-methylmethionine	Dnmmet
	D-N-methylornithine	Dnmorn	N-methylcyclopentylalanine	Nmcpen

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	N-methylglycine	Nala	D-N-methylphenylalanine	Dnmphe
	N-methylaminoisobutyrate	Nmaib	D-N-methylproline	Dnmpro
	N-(1-methylpropyl)glycine	Nile	D-N-methylserine	Dnmser
	N-(2-methylpropyl)glycine	Nleu	D-N-methylthreonine	Dnmthr
5	D-N-methyltryptophan	Dnmtrp	N-(1-methylethyl)glycine	Nval
	D-N-methyltyrosine	Dnmtyr	N-methyl- <i>n</i> -naphthylalanine	Nmanap
	D-N-methylvaline	Dnmval	N-methylpenicillamine	Nmpen
	γ -aminobutyric acid	Gabu	N-(<i>p</i> -hydroxyphenyl)glycine	Nhtyr
	L- <i>t</i> -butylglycine	Tbug	N-(thiomethyl)glycine	Ncys
10	L-ethylglycine	Etg	penicillamine	Pen
	L-homophenylalanine	Hphe	L- α -methylalanine	Mala
	L- α -methylarginine	Marg	L- α -methylassparagine	Masn
	L- α -methylasspartate	Masp	L- α -methyl- <i>t</i> -butylglycine	Mtbug
	L- α -methylcysteine	Mcys	L-methylethylglycine	Metg
15	L- α -methylglutamine	Mgln	L- α -methylglutamate	Mglu
	L- α -methylhistidine	Mhis	L- α -methylhomophenylalanine	Mhphe
	L- α -methylisoleucine	Mile	N-(2-methylthioethyl)glycine	Nmet
	L- α -methyllleucine	Mleu	L- α -methyllysine	Mlys
	L- α -methylmethionine	Mmet	L- α -methylnorleucine	Mnle
20	L- α -methylnorvaline	Mnva	L- α -methylornithine	Morn
	L- α -methylphenylalanine	Mphe	L- α -methylproline	Mpro
	L- α -methylserine	Mser	L- α -methylthreonine	Mthr
	L- α -methyltryptophan	Mtrp	L- α -methyltyrosine	Mtyr
	L- α -methylvaline	Mval	L-N-methylhomophenylalanine	Nmhpe
25	N-(N-(2,2-diphenylethyl)carbamylmethyl)glycine	Nnbhm	N-(N-(3,3-diphenylpropyl)carbamylmethyl)glycine	Nnbhe
	1-carboxy-1-(2,2-diphenyl-Nmbe ethylamino)cyclopropane			

30

Crosslinkers can be used, for example, to stabilise 3D conformations, using homo-bifunctional crosslinkers such as the bifunctional imido esters having (CH₂)_n spacer groups with n=1 to n=6, glutaraldehyde, N-hydroxysuccinimide esters and hetero-bifunctional

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reagents which usually contain an amino-reactive moiety such as N-hydroxysuccinimide and another group specific-reactive moiety.

Still without limiting the present invention to any one theory or mode of action, it is
5 thought that the human wild-type sphingosine kinase protein region defined by amino acid residues 16-153 comprises all or part of an ATP binding site. Accordingly, it is thought that by blocking the ATP binding site, the subject sphingosine kinase is rendered catalytically inactive in terms of its capacity to phosphorylate sphingosine to sphingosine-
1-phosphate. The phrase "functionally equivalent region" should therefore be understood
10 as a reference to any region of a sphingosine kinase amino acid sequence which exhibits at least one of the function activities attributable to the region defined by amino acid residue numbers 16-153.

Accordingly, in another aspect the present invention is directed to a sphingosine kinase
15 variant comprising a mutation in an ATP binding site region or functionally equivalent region wherein said variant exhibits ablated or reduced catalytic activity relative to wild-type sphingosine kinase or a derivative, homologue, analogue, chemical equivalent or mimetic of said sphingosine kinase variant .

20 Preferably, said sphingosine kinase is a human sphingosine kinase.

Still more preferably, said mutation is a substitution, deletion and/or addition of one or more amino acids in the region defined by amino acid residues 16-153, more preferably
70-90 and still more preferably 79-84.

25

In a most preferred embodiment, said mutation comprises one or more of the amino acid substitutions selected from the following list:

- (i) G82D
- (ii) G82A
- 30 (iii) G26D
- (iv) S79D

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- (v) G80D
- (vi) K103A
- (vii) G111D
- (viii) G113D
- 5 (ix) G26A
- (x) K27A
- (xi) K29A
- (xii) S79A
- (xiii) G80A
- 10 (xiv) K103R
- (xv) G111A

To the extent that the present invention relates to sphingosine kinase variants comprising one or more amino acid additions, substitutions and/or deletions, it should also be
15 understood to extend to nucleic acid molecules encoding said variants.

Accordingly, another aspect of the present invention is directed to an isolated nucleic acid molecule selected from the list consisting of:

- 20 (i) An isolated nucleic acid molecule or derivative or equivalent thereof comprising a nucleotide sequence encoding or complementary to a sequence encoding a sphingosine kinase variant or derivative, homologue, analogue, chemical equivalent or mimetic of said variant which variant comprises a mutation in a region defined by amino acid 16-153 or functionally equivalent region wherein said variant
25 exhibits ablated or reduced catalytic activity relative to wild-type sphingosine kinase.
- (ii) An isolated nucleic acid molecule or derivative or equivalent thereof comprising a nucleotide sequence encoding or complementary to a sequence encoding a human
30 sphingosine kinase variant or derivative, homologue, analogue, chemical equivalent or mimetic of said variant which variant comprises a mutation in a region defined

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by amino acid 16-153 or functionally equivalent region wherein said variant exhibits ablated or reduced catalytic activity relative to wild-type human sphingosine kinase.

- 5 (iii) An isolated nucleic acid molecule or derivative or equivalent thereof comprising a nucleotide sequence encoding or complementary to a sequence encoding a human sphingosine kinase variant or derivative, homologue, analogue, chemical equivalent or mimetic of said variant, which variant comprises an amino acid sequence with a single or multiple multiple amino acid substitution, addition and/or deletion in a
10 region defined by amino acid 16-153 or functionally equivalent region wherein said variant exhibits ablated or reduced catalytic activity relative to wild-type sphingosine kinase.
- (iv) An isolated nucleic acid molecule or derivative or equivalent thereof comprising a
15 nucleotide sequence encoding or complementary to a sequence encoding a human sphingosine kinase variant or derivative, homologue, analogue, chemical equivalent or mimetic of said variant, which variant comprises an amino acid sequence with a single or multiple amino acid substitution, addition and/or deletion in a region defined by amino acid 70-90 or functionally equivalent region wherein said variant
20 exhibits ablated or reduced catalytic activity relative to wild-type sphingosine kinase.
- (v) An isolated nucleic acid molecule or derivative or equivalent thereof comprising a
25 nucleotide sequence encoding or complementary to a sequence encoding a human sphingosine kinase variant or derivative, homologue, analogue, chemical equivalent or mimetic of said variant, which variant comprises an amino acid sequence with a single or multiple multiple amino acid substitution, addition and/or deletion in a region defined by amino acid 79-84 or functionally equivalent region wherein said variant exhibits ablated or reduced catalytic activity relative to wild-type
30 sphingosine kinase.

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- (vi) An isolated nucleic acid molecule or derivative or equivalent thereof comprising a nucleotide sequence encoding or complementary to a sequence encoding a sphingosine kinase variant or a derivative, homologue, analogue, chemical equivalent or mimetic of said variant comprising one or more of the amino acid substitutions selected from the following list:
- 5
- (a) G82D
 - (b) G82A
 - (c) G26D
 - (d) S79D
 - 10 (e) G80D
 - (f) K103A
 - (g) G111D
 - (h) G113D
 - (i) G26A
 - 15 (j) K27A
 - (k) K29A
 - (l) S79A
 - (m) G80A
 - (n) K103R
 - 20 (o) G111A
- (vii) An isolated nucleic acid molecule or derivative or analogue thereof comprising a nucleotide sequence encoding or complementary to a sequence encoding a sphingosine kinase variant or derivative, homologue, analogue, chemical equivalent or mimetic of said variant which variant comprises a mutation in an ATP binding site region or functionally equivalent region wherein said variant exhibits ablated or reduced catalytic activity relative to wild-type sphingosine kinase.
- 25

The nucleic acid molecule of the subject invention may be ligated to an expression vector capable of expression in a prokaryotic cell (eg. *E. Coli*) or a eukaryotic cell (eg. yeast cells, fungal cells, insect cells, mammalian cells or plant cells). The nucleic acid molecule may

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be ligated or fused or otherwise associated with a nucleic acid molecule encoding another entity such as, for example, a signal peptide. It may also comprise additional nucleotide sequence information fused, linked or otherwise associated with it either at the 3' or 5' terminal portions or at both the 3' and 5' terminal portions. The nucleic acid molecule may
5 also be part of a vector, such as an expression vector. The latter embodiment facilitates production of recombinant forms of the variant sphingosine kinase encompassed by the present invention.

The variant sphingosine kinase molecule of the present invention may be derived from
10 natural or recombinant sources or may be chemically synthesised. Methods for producing these molecules would be well known to those skilled in the art.

In addition to facilitating the synthesis of sphingosine kinase variants, *per se*, identification of the mechanism of functioning of the sphingosine kinase variants of the present invention
15 permits the design of methodology for ablating or decreasing the catalytic activity of wild-type sphingosine kinase proteins. For example, contacting wild-type sphingosine kinase proteins with an agent which bind to or otherwise associate with the region defined by amino acids 16-153 or functionally equivalent region is possible to effectively convert a wild-type sphingosine kinase protein to a catalytically inactive variant. In another aspect,
20 said agent could bind to or otherwise associate with the sphingosine kinase ATP binding site.

Without limiting the present invention in any way, it is thought that baseline sphingosine kinase activity is a constitutive property of all wild-type sphingosine kinase proteins. In
25 order to activate this molecule, though, one has to increase the V_{max} of the enzyme (eg., by transporting more of the enzyme to the location at which it is required or by altering its function post-translationally). This requires the functioning of another molecule or class of molecules (such as a protein or lipid) to associate with the subject sphingosine kinase, these molecules hypothetically being termed FOSK(s) (Friends of Sphingosine Kinase).
30 Reference to "FOSK" should be understood to include reference to sphingosine kinase interacting molecules (SKIMS). The G82D sphingosine kinase variant, for example, is

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thought to bind a FOSK molecule thereby preventing it from activating wild-type sphingosine kinase.

Accordingly, the present invention provides not only sphingosine kinase variants, *per se*, as potential drugs, but provides a mechanism for further drug development. For example,
5 a small molecule that binds to the region which is the subject of mutation in the sphingosine kinase variant molecule would be expected to convert a wild-type sphingosine kinase protein to an inhibitor that not only loses its baseline sphingosine kinase activity but also causes its conversion into a dominant negative sphingosine kinase variant, ie., both levels of sphingosine kinase activity are eliminated. Alternatively, screening for an agent
10 which prevents interaction of the wild-type sphingosine kinase with a FOSK would reproduce the dominant negative phenotype while not disturbing baseline sphingosine kinase activity but only inhibiting activation. This is a potentially desirable scenario where some sphingosine kinase activity is required for cellular survival.

15 Modulation of the activity between sphingosine kinase may therefore be achieved by any one of a number of techniques including, but not limited to:

- (i) introducing into a cell a proteinaceous or non-proteinaceous molecule which antagonises the interaction between a FOSK and sphingosine kinase.
20
- (ii) introducing into a cell a proteinaceous or non-proteinaceous molecule which interacts with at least part of the region of wild type sphingosine kinase which is the subject mutation in the variants described herein.

25 Reference to "agent" should therefore be understood as a reference to any proteinaceous or non-proteinaceous molecule which modulates the interaction of sphingosine kinase with a FOSK or interacts with at least part of the region defined by amino acids 16-153 of sphingosine kinase and includes, for example, the molecules detailed in points (i) – (ii), above. The subject agent may be linked, bound or otherwise associated with any
30 proteinaceous or non-proteinaceous molecule. For example, it may be associated with a molecule which permits its targeting to a localised region.

- 30 -

Accordingly, by administering said agent intracellularly, endogenously produced wild-type sphingosine kinase could be both inactivated in terms of its baseline activity and induced to function as a dominant negative sphingosine kinase molecule wherein the agent-

5 associated wild-type sphingosine kinase functions to ablate or decrease activation of other non-agent associated wild-type sphingosine kinase molecules. Alternatively, since in some instances it is necessary to maintain intracellular baseline sphingosine kinase activity, the agent could be associated with wild-type sphingosine kinase extracellularly and the agent-sphingosine kinase complex could then be administered intracellularly. These complexes

10 (being a variant sphingosine kinase within the context of the present invention) would then act to ablate or decrease wild-type sphingosine kinase activation without significantly modulating baseline activity of endogenously produced wild-type sphingosine kinase proteins. The design and generation of these molecules provides a unique and previously unavailable mechanism for modulating activity of the sphingosine kinase signalling path.

15 Specifically, whereas previously utilised chemical inhibitors such as N,N'-dimethylsphingosine totally eliminate sphingosine kinase functioning, the variants of the present invention can be administered such as to only reduce or eliminate activation of wild-type sphingosine kinase without disturbing baseline functioning.

20 The subject agent may be any proteinaceous or non-proteinaceous molecule derived from natural, recombinant or synthetic sources including fusion proteins or following, for example, natural product screening and which achieves the object of the present invention. Synthetic sources of said agent include for example chemically synthesised molecules. In other examples, phage display libraries can be screened for peptides while chemical

25 libraries can be screened for existing small molecules. Rational drug design/structure based design can be achieved by performing crystallisation, further analysing the ATP binding site and fitting molecules into that site by design.

By way of example, diversity libraries, such as random combinatorial peptide or

30 nonpeptide libraries can be screened. Many publically or commercially available libraries

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can be used such as chemically synthesized libraries, recombinant (e.g., phage display libraries) and *in vitro* translation-based libraries.

Examples of chemically synthesized libraries are described in Fodor *et al.*, (1991); Houghten *et al.*, (1991); Lam *et al.*, (1991); Medynski., (1994); Gallop *et al.*, (1994);
5 Ohlmeyer *et al.*, (1993); Erb *et al.*, (1994); Houghten *et al.*, (1992); Jayawickreme *et al.*, (1994); Salmon *et al.*, (1993); International Patent Publication No. WO 93/20242; and Brenner and Lerner., (1992).

Examples of phage display libraries are described by Scott and Smith., (1990); Devlin *et al.*, (1990); Christian R.B *et al.*, (1992); Lenstra., (1992); Kay *et al.*, (1993) and
10 International Patent Publication No. WO 94/18318.

In vitro translation-based libraries include but are not limited to those described in Mattheakis *et al.*, (1994).

15 Screening the libraries can be accomplished by any of a variety of commonly known methods. See Parmley and Smith., (1989); Scott and Smith., (1990); Fowlkes *et al.*, (1992); Oldenburg *et al.*, (1992); Yu *et al.*, (1994); Staudt *et al.*, (1988); Bock *et al.*, (1992); Tuerk *et al.*, (1992); Ellington *et al.*, (1992); U.S. Patent No. 5,096,815, U.S. Patent No. 5,223,409 and U.S. Patent No. 5,198,346; Rebar and Pabo., (1993); and
20 International Patent Publication No. WO 94/18318.

The present invention should therefore also be understood to extend to a method of screening for agents which modulate the interaction between sphingosine kinase and a
25 FOSK molecule. This could be achieved, for example, by utilising cell based assays which can monitor sphingosine kinase activation. Accordingly, the present invention provides a mechanism of screening for agents which utilise one of a variety of methods of inhibiting sphingosine kinase. For example, agents which totally ablate both levels of sphingosine kinase activity can be screened for in addition to molecules which only inhibit sphingosine
30 kinase activation (for example by inhibiting the interaction between sphingosine kinase and a FOSK).

Screening for the modulatory agents hereinbefore defined can be achieved by any one of several suitable methods including, but in no way limited to, contacting a cell comprising sphingosine kinase (separately or together with FOSK) with an agent and screening for
5 the modulation of sphingosine kinase/FOSK functional activity or modulation of the activity or expression of a downstream sphingosine kinase or FOSK cellular target. Detecting such modulation can be achieved utilising techniques such as Western blotting, electrophoretic mobility shift assays and/or the readout of reporters of sphingosine kinase or FOSK activity such as luciferases, CAT and the like.

10

It should be understood that the sphingosine kinase or FOSK protein may be naturally occurring in the cell which is the subject of testing or the genes encoding them may have been transfected into a host cell for the purpose of testing. Further, the naturally occurring or transfected gene may be constitutively expressed - thereby providing a model useful for,
15 *inter alia*, screening for agents which down-regulate sphingosine kinase FOSK interactivity or the gene may require activation - thereby providing a model useful for, *inter alia*, screening for agents which modulate sphingosine kinase/FOSK interactivity under certain stimulatory conditions. Further, to the extent that a sphingosine kinase nucleic acid molecule is transfected into a cell, that molecule may comprise the entire sphingosine
20 kinase gene or it may merely comprise a portion of the gene such as the FOSK binding portion.

In another example, the subject of detection could be a downstream sphingosine kinase regulatory target, rather than sphingosine kinase itself. Yet another example includes
25 sphingosine kinase binding sites ligated to a minimal reporter. For example, modulation of sphingosine kinase/FOSK interactivity can be detected by screening for the modulation of the downstream signalling components of a TNF stimulated cell. This is an example of a system where modulation of the molecules which sphingosine kinase and FOSK regulate the activity of, are monitored.

30

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Accordingly, another aspect of the present invention provides a method for detecting an agent capable of modulating the interaction of FOSK with sphingosine kinase or its functional equivalent or derivative thereof said method comprising contacting a cell or extract thereof containing said sphingosine kinase and FOSK or its functional equivalent or
5 derivative with a putative agent and detecting an altered expression phenotype associated with said interaction.

Reference to "sphingosine kinase" and "FOSK" should be understood as a reference to either the sphingosine kinase or FOSK expression product or to a portion or fragment of
10 the sphingosine kinase or FOSK molecule, such as the FOSK region defined by amino acids 16-153 of the sphingosine kinase protein. In this regard, the sphingosine kinase or FOSK expression product is expressed in a cell. The cell may be a host cell which has been transfected with the sphingosine kinase or FOSK nucleic acid molecule or it may be a cell which naturally contains the sphingosine kinase gene. Reference to "extract thereof"
15 should be understood as a reference to a cell free transcription system.

Reference to detecting an "altered expression phenotype associated with said interaction" should be understood as the detection of cellular changes associated with modulation of the interaction of sphingosine kinase with FOSK. These may be detectable, for example, as
20 intracellular changes or changes observable extracellularly. For example, this includes, but is not limited to, detecting changes in downstream product levels or activities.

In yet another aspect the present invention provides a method for detecting an agent capable of binding or otherwise associating with the sphingosine kinase region defined by
25 amino acids 16-153 or functional equivalent or derivative thereof said method comprising contacting a cell containing said amino acid region or functional equivalent or derivative thereof with a putative agent and detecting an altered expression phenotype associated with modulation of the function of sphingosine kinase or its functional equivalent or derivative.
30 Preferably, said region is defined by amino acids 70-90 and even more preferably 79-84.

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Reference to "sphingosine kinase binding site" should be understood as a reference to the sphingosine kinase region defined by amino acids 16-153, preferably 70-90 and even more preferably 79-84.

- 5 In addition to screening for agents which modulate the interaction of FOSK and sphingosine kinase utilising function based assays of the type described above, the identification of the functionally active region of sphingosine kinase also facilitates the screening, analysis, rational design and/or modification of agents for modulating either the interaction of FOSK and sphingosine kinase or the activity of sphingosine kinase based on
10 analysis of the physical interaction of a putative agent or lead compound with the subject region.

Specifically, knowledge of the nature and location of this site now facilitates analysis of the tertiary structure of sphingosine kinase, in terms of the structure of the binding site, by
15 techniques such as X-ray crystallography.

Accordingly, another aspect of the present invention is directed to a method for analysing, designing and/or modifying an agent capable of interacting with the sphingosine kinase region defined by amino acids 16-153 or derivative thereof and modulating at least one
20 functional activity associated with said sphingosine kinase said method comprising contacting said sphingosine kinase or derivative thereof with a putative agent and assessing the degree of interactive complementarity of said agent with said binding site.

Preferably, said region is defined by amino acids 70-90 and even more preferably 79-84.
25

It should be understood that the sphingosine kinase which is contacted with the putative agent for evaluation of interactive complementarity may be recombinantly produced. However, it should also be understood that the subject sphingosine kinase may take the form of an image based on the binding site structure which has been elucidated, such as an
30 electron density map, molecular models (including, but not limited to, stick, ball and stick, space filling or surface representation models) or other digital or non-digital surface

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representation models or image, which facilitates the analysis of sphingosine kinase site: agent interactions utilising techniques and software which would be known to those of skill in the art. For example, interaction analyses can be performed utilising techniques such as Biacore real-time analysis of on and off-rates and dissociation constants for binding of
5 ligands (Gardsvoll *et al*, 1999; Hoyer-Hansen *et al*, 1997; Ploug, 1998; Ploug *et al*, 1994; 1995; 1998) and NMR perturbation studies (Stephens *et al*, 1992).

Reference to "assessing the degree of interactive complementarity" of an agent with the subject sphingosine kinase binding site should be understood as a reference to elucidating
10 any feature of interest including, but not limited to, the nature and/or degree of interaction between the subject sphingosine kinase binding site and an agent of interest. As detailed above, any suitable technique can be utilised. Such techniques would be known to the person of skill in the art and can be utilised in this regard. In terms of the nature of the subject interaction, it may be desirable to assess the types of interactive mechanisms which
15 occur between specific residues of any given agent and those of the sphingosine kinase binding site (for example, peptide bonding or formation of hydrogen bonds, ionic bonds, van der Waals forces, etc.) and/or their relative strengths. It may also be desirable to assess the degree of interaction which occurs between an agent of interest and the subject sphingosine kinase binding site. For example, by analysing the location of actual sites of
20 interaction between the subject agent and sphingosine kinase binding site it is possible to determine the quality of fit of the agent into this region of the sphingosine kinase binding site and the relative strength and stability of that binding interaction. For example, if it is the object that sphingosine kinase binding site functioning be blocked, an agent which interacts with the sphingosine kinase binding site such that it blocks or otherwise hinders
25 (for example, sterically hinders or chemically or electrostatically repels) FOSK interaction or down-regulates sphingosine kinase activity will be sought. The form of association which is required in relation to modulating sphingosine kinase functioning may not involve the formation of any chemical interactive bonding mechanism, as this is traditionally understood, but may involve a non-bonding mechanism such as the proximal location of a
30 region of the agent relative to the subject binding region of the sphingosine kinase binding site, for example, to effect steric hindrance with respect to the binding of an activating

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molecule. Where the interaction takes the form of hindrance or the creation of other repulsive forces, this should nevertheless be understood as a form of "interaction" despite the lack of formation of any of the traditional forms of bonding mechanisms.

5 It should also be understood that the sphingosine kinase binding site which is utilised either in a physical form or as an image, as hereinbefore discussed, to assess the interactive complementarity of a putative agent may be a naturally occurring form of the sphingosine kinase binding site or it may be a derivative, homologue, analogue, mutant, fragment or equivalent thereof. The derivative, homologue, analogue, mutant, fragment or equivalent
10 thereof may take either a physical or non-physical (such as an image) form.

The determination of sphingosine kinase binding regions facilitates determination of the three dimensional structure of the sphingosine kinase binding site and the identification and/or rational modification and design of agents which can be used to modulate FOSK
15 binding or sphingosine kinase functioning.

Without limiting the application of the present invention in any way, the method of the present invention facilitates the analysis, design and/or modification of agents capable of interacting with the sphingosine kinase site defined by amino acids 16-153. In this regard,
20 reference to "analysis, design and/or modification" of an agent should be understood in its broadest sense to include:

(i) Randomly screening (for example, utilising routine high-throughput screening technology) to identify agents which exhibit some modulatory capacity with
25 respect to sphingosine kinase functional activity and/or FOSK binding and then analysing the precise nature and magnitude of the agent's modulatory capacity utilising the method of this aspect of the present invention. In this regard, existing crystals could be soaked with said agents or co-crystallisation could be performed. A combination of modelling and synthetic modification of the local compound
30 together with mutagenesis of the sphingosine kinase binding site could then be performed for example. In screening for agents which may modulate activity,

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standard methods of phage display and also combinatorial chemistry may be utilised (Goodson *et al.*, 1994; Terrett., 2000). Such interaction studies can also be furthered utilising techniques such as the Biacore analysis and NMR perturbation studies. Such agents are often commonly referred to as "lead" agents in terms of the random screening of proteinaceous or non-proteinaceous molecules for their capacity to function either agonistically or antagonistically. Further, for example, binding affinity and specificity could be enhanced by modifying lead agents to maximise interactions with the sphingosine kinase binding site. Such analyses would facilitate the selection of agents which are the most suitable for a given purpose. In this way, the selection step is based not only on *in vitro* data but also on a technical analysis of sites of agent: sphingosine kinase interaction in terms of their frequency, stability and suitability for a given purpose. For example, such analysis may reveal that what appears to be an acceptable *in vitro* activity in respect of a randomly identified agent is in fact induced by a highly unstable interaction due to the presence of proximally located agent: sphingosine kinase sites which exhibit significant repulsive forces thereby de-stabilising the overall interaction between the agent and the sphingosine kinase. This would then facilitate the selection of another prospective lead compound, exhibiting an equivalent degree of *in vitro* activity, but which agent does not, upon further analysis, involve the existence of such de-stabilising repulsive forces.

Screening for the modulatory agents herein defined can be achieved by any one of several suitable methods, including *in silico* methods, which would be well known to those of skill in the art and which are, for example, routinely used to randomly screen proteinaceous and non-proteinaceous molecules for the purpose of identifying lead compounds.

These methods provide a mechanism for performing high throughput screening of putative modulatory agents such as the proteinaceous or non-proteinaceous agents comprising synthetic, recombinant, chemical and natural libraries.

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- 5 (ii) The candidate or lead agent (for example, the agent identified in accordance with the methodology described in relation to point (i)) could be modified in order to maximise desired interactions (for example, binding affinity to specificity) with the sphingosine kinase and to minimise undesirable interactions (such as repulsive or otherwise de-stabilising interactions).

10 Methods of modification of a candidate or lead agent in accordance with the purpose as defined herein would be well known to those of skill in the art. For example, a molecular replacement program such as Amore (Navaza, 1994) may be utilised in this regard. The method of the present invention also facilitates the mutagenesis of known signal inducing agents in order to ablate or improve signalling activity.

- 15 (iii) In addition to analysing fit and/or structurally modifying existing molecules, the method of the present invention also facilitates the rational design and synthesis of an agent, such as an agonistic or antagonistic agent, based on theoretically modelling an agent exhibiting the desired sphingosine kinase binding site interactive structural features followed by the synthesis and testing of the subject agent.

20

It should be understood that any one or more of applications (i) – (iii) above, may be utilised in identifying a particular agent.

25 In a related aspect, the present invention should be understood to extend to the agents identified utilising any of the methods hereinbefore defined. In this regard, reference to an agent should be understood as a reference to any proteinaceous or non-proteinaceous molecule which modulates at least one sphingosine kinase functional activity.

30 Without limiting the theory or mode of action of the present invention, sphingosine kinase is a key regulatory enzyme in the activity of the sphingosine kinase signalling pathway.

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By "sphingosine kinase signalling pathway" is meant a signalling pathway which utilises one or both of sphingosine kinase and/or sphingosine-1-phosphate. It is thought that a sphingosine kinase signalling pathway cascade may take the form of:

- 5 (i) the generation of ceramide from sphingomyelin via *S.Mase* activity, said ceramide being converted to sphingosine;
- (ii) sphingosine-1-phosphate generation by stimulation of sphingosine kinase; and
- 10 (iii) the activation of MEK/ERK and nuclear translocation of NF- κ B downstream from Sphingosine-1-phosphate generation.

The sphingosine kinase signalling pathway is known to regulate cellular activities such as those which lead to inflammation, apoptosis and cell proliferation. For example, upregulation of the production of inflammatory mediators such as cytokines, chemokines, eNOS and upregulation of adhesion molecule expression. Said upregulation may be induced by a number of stimuli including, for example, inflammatory cytokines such as tumour necrosis factor- α (TNF- α) and interleukin-1 (IL-1), endotoxin, oxidised or modified lipids, radiation or tissue injury.

The generation of variant sphingosine kinase molecules now provides additional molecules for use in the prophylactic and therapeutic treatment of diseases characterised by unwanted cellular activity, which activity is either directly or indirectly modulated via activity of the sphingosine kinase signalling pathway. Examples of diseases involving unwanted sphingosine kinase regulated cellular activity include inflammatory conditions (eg., rheumatoid arthritis, inflammatory bowel disease), neoplastic conditions (eg., solid cancers), asthma, atherosclerosis, meningitis, multiple sclerosis and septic shock. The variants of the present invention may also facilitate the provision of chronic treatment in relation to disease conditions such as atherosclerosis, osteoarthritis and other degenerative diseases in which inflammation plays a role.

- 40 -

Accordingly, the present invention contemplates therapeutic and prophylactic uses of variant sphingosine kinase molecules for the regulation of cellular functional activity, such as for example, regulation of inflammation. In this regard, the variant molecules which may be used in therapy and prophylaxis include mutated sphingosine kinase expression
5 product, nucleic acid molecules encoding mutated sphingosine kinase expression product, sphingosine kinase-agent complexes as hereinbefore defined or an agent, *per se*, which is proposed to be administered to a subject for the purpose of its intracellular complexation with wild-type sphingosine kinase for the purpose of converting a wild-type molecule to a variant sphingosine kinase molecule. For ease of reference and in accordance with the
10 definitions provided earlier, it should be understood that the phrase "sphingosine kinase variant" includes reference to mutated sphingosine kinase proteins, nucleic acid molecules encoding said proteins and sphingosine kinase-agent complexes while reference to "agent" is intended to refer to an agent which, when contacted with a sphingosine kinase protein (such as a wild-type protein) will render the protein a variant within the context of the
15 present invention.

Accordingly, another aspect of the present invention contemplates a method of modulating cellular functional activity in a mammal said method comprising administering to said mammal an effective amount of a sphingosine kinase variant or agent as hereinbefore
20 defined for a time and under conditions sufficient to inhibit, reduce or otherwise down-regulate at least one functional activity of wild-type sphingosine kinase.

Preferably said functional activity is down-regulation of wild-type sphingosine kinase baseline activity and/or prevention of wild-type sphingosine kinase activation.
25

Reference to "modulating cellular functional activity" is a reference to up-regulating, down-regulating or otherwise altering any one or more of the activities which a cell is capable of performing such as, but not limited to, one or more of chemokine production, cytokine production, nitric oxide synthetase, adhesion molecule expression and production
30 of other inflammatory modulators.

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Administration of the variant sphingosine kinase or agent, in the form of a pharmaceutical composition, may be performed by any convenient means. Variant sphingosine kinase or agent of the pharmaceutical composition are contemplated to exhibit therapeutic activity when administered in an amount which depends on the particular case. The variation
5 depends, for example, on the human or animal and the sphingosine kinase or agent chosen. A broad range of doses may be applicable. Considering a patient, for example, from about 0.1 mg to about 1 mg of sphingosine kinase or agent may be administered per kilogram of body weight per day. Dosage regimes may be adjusted to provide the optimum therapeutic response. For example, several divided doses may be administered daily, weekly, monthly
10 or other suitable time intervals or the dose may be proportionally reduced as indicated by the exigencies of the situation. The variant sphingosine kinase or agent may be administered in a convenient manner such as by the oral, intravenous (where water soluble), intranasal, intraperitoneal, intramuscular, subcutaneous, intradermal or suppository routes or implanting (e.g. using slow release molecules). With particular
15 reference to use of variant sphingosine kinase or agent, these molecules may be administered in the form of pharmaceutically acceptable nontoxic salts, such as acid addition salts or metal complexes, e.g. with zinc, iron or the like (which are considered as salts for purposes of this application). Illustrative of such acid addition salts are hydrochloride, hydrobromide, sulphate, phosphate, maleate, acetate, citrate, benzoate,
20 succinate, malate, ascorbate, tartrate and the like. If the active ingredient is to be administered in tablet form, the tablet may contain a binder such as tragacanth, corn starch or gelatin; a disintegrating agent, such as alginic acid; and a lubricant, such as magnesium stearate.

25 A further aspect of the present invention relates to the use of the invention in relation to mammalian disease conditions. For example, the present invention is particularly useful, but in no way limited to, use in therapeutically or prophylactically treating inflammatory diseases, neoplastic conditions and degenerative diseases.

30 Accordingly, another aspect of the present invention relates to the treatment and/or prophylaxis of a condition in a mammal, which condition is characterised by aberrant,

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unwanted or otherwise inappropriate cellular activity, said method comprising administering to said mammal an effective amount of a sphingosine kinase variant or agent as hereinbefore defined for a time and under conditions sufficient to inhibit, reduce or otherwise down-regulate at least one functional activity of wild-type sphingosine kinase
5 wherein said down-regulation results in modulation of cellular functional activity.

Preferably said functional activity is down-regulation of baseline wild-type sphingosine kinase activity and/or prevention of wild-type sphingosine kinase activation.

10 Reference to "aberrant, unwanted or otherwise inappropriate" cellular activity should be understood as a reference to overactive cellular activity, underactive cellular activity or physiologically normal cellular activity which is inappropriate or otherwise unwanted.

The subject of the treatment or prophylaxis is generally a mammal such as but not limited
15 to human, primate, livestock animal (eg. sheep, cow, horse, donkey, pig), companion animal (eg. dog, cat), laboratory test animal (eg. mouse, rabbit, rat, guinea pig hamster), captive wild animal (eg. fox, deer). Preferably the mammal is human or primate. Most preferably the mammal is a human.

20 Reference herein to "treatment" and "prophylaxis" is to be considered in its broadest context. The term "treatment" does not necessarily imply that a mammal is treated until total recovery. Similarly, "prophylaxis" does not necessarily mean that the subject will not eventually contract a disease condition. Accordingly, treatment and prophylaxis including amelioration of the symptoms of a particular condition or preventing or otherwise reducing
25 the risk of developing a particular condition. The term "prophylaxis" may be considered as reducing the severity or onset of a particular condition. "Treatment" may also reduce the severity of an existing condition.

An "effective amount" means an amount necessary at least partly to attain the desired
30 immune response, or to delay the onset or inhibit progression or halt altogether, the onset or progression of a particular condition of the individual to be treated, the taxonomic group

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of individual to be treated, the degree of protection desired, the formulation of the vaccine, the assessment of the medical situation, and other relevant factors. It is expected that the amount will fall in a relatively broad range that can be determined through routine trials.

- 5 A further aspect of the present invention relates to the use of a sphingosine kinase variant or agent as hereinbefore defined in the manufacture of a medicament for the modulation of cellular functional activity.

Another aspect of the present invention relates to a sphingosine kinase variant or agent as
10 hereinbefore defined for use in modulating cellular functional activity.

In a related aspect of the present invention, the mammal undergoing treatment may be a human or an animal in need of therapeutic or prophylactic treatment.

- 15 In accordance with these methods, the molecules defined in accordance with the present invention may be coadministered with one or more other compounds or molecules. By "coadministered" is meant simultaneous administration in the same formulation or in two different formulations via the same or different routes or sequential administration by the same or different routes. By "sequential" administration is meant a time difference of from
20 seconds, minutes, hours or days between the administration of the two types of molecules. These molecules may be administered in any order.

- In yet another further aspect the present invention contemplates a pharmaceutical composition comprising a sphingosine kinase variant or agent as hereinbefore defined
25 together with one or more pharmaceutically acceptable carriers and/or diluents. The sphingosine kinase variant and agent are referred to as the active ingredients.

- The pharmaceutical forms suitable for injectable use include sterile aqueous solutions (where water soluble) and sterile powders for the extemporaneous preparation of sterile
30 injectable solutions or dispersion. In all cases the form must be sterile and must be fluid to the extent that easy syringability exists. It must be stable under the conditions of

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manufacture and storage and must be preserved against the contaminating action of microorganisms such as bacteria and fungi. The carrier can be a solvent or dispersion medium containing, for example, water, ethanol, polyol (for example, glycerol, propylene glycol and liquid polyethylene glycol, and the like), suitable mixtures thereof, and
5 vegetable oils. The proper fluidity can be maintained, for example, by the use of a coating such as lecithin, by the maintenance of the required particle size in the case of dispersion and by the use of surfactants. The preventions of the action of microorganisms can be brought about by various antibacterial and antifungal agents, for example, parabens, chlorobutanol, phenol, sorbic acid, thimerosal and the like. In many cases, it will be
10 preferable to include isotonic agents, for example, sugars or sodium chloride. Prolonged absorption of the injectable compositions can be brought about by the use in the compositions of agents delaying absorption, for example, aluminum monostearate and gelatin.

Sterile injectable solutions are prepared by incorporating the active compounds in the
15 required amount in the appropriate solvent with various of the other ingredients enumerated above, as required, followed by filtered sterilization. Generally, dispersions are prepared by incorporating the various sterilized active ingredient into a sterile vehicle which contains the basic dispersion medium and the required other ingredients from those enumerated above. In the case of sterile powders for the preparation of sterile injectable
20 solutions, the preferred methods of preparation are vacuum drying and the freeze-drying technique which yield a powder of the active ingredient plus any additional desired ingredient from previously sterile-filtered solution thereof.

When the active ingredients are suitably protected they may be orally administered, for
25 example, with an inert diluent or with an assimilable edible carrier, or they may be enclosed in hard or soft shell gelatin capsule, or they may be compressed into tablets, or they may be incorporated directly with the food of the diet. For oral therapeutic administration, the active compound may be incorporated with excipients and used in the form of ingestible tablets, buccal tablets, troches, capsules, elixirs, suspensions, syrups,
30 wafers, and the like. Such compositions and preparations should contain at least 1% by weight of active compound. The percentage of the compositions and preparations may, of

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course, be varied and may conveniently be between about 5 to about 80% of the weight of the unit. The amount of active compound in such therapeutically useful compositions in such that a suitable dosage will be obtained. Preferred compositions or preparations according to the present invention are prepared so that an oral dosage unit form contains
5 between about 0.1 μ g and 2000 mg of active compound.

The tablets, troches, pills, capsules and the like may also contain the following: A binder such as gum tragacanth, acacia, corn starch or gelatin; excipients such as dicalcium phosphate; a disintegrating agent such as corn starch, potato starch, alginic acid and the
10 like; a lubricant such as magnesium stearate; and a sweetening agent such a sucrose, lactose or saccharin may be added or a flavouring agent such as peppermint, oil of wintergreen, or cherry flavouring. When the dosage unit form is a capsule, it may contain, in addition to materials of the above type, a liquid carrier. Various other materials may be present as coatings or to otherwise modify the physical form of the dosage unit. For
15 instance, tablets, pills, or capsules may be coated with shellac, sugar or both. A syrup or elixir may contain the active compound, sucrose as a sweetening agent, methyl and propylparabens as preservatives, a dye and flavouring such as cherry or orange flavour. Of course, any material used in preparing any dosage unit form should be pharmaceutically pure and substantially non-toxic in the amounts employed. In addition, the active
20 compound may be incorporated into sustained-release preparations and formulations.

Pharmaceutically acceptable carriers and/or diluents include any and all solvents, dispersion media, coatings, antibacterial and antifungal agents, isotonic and absorption delaying agents and the like. The use of such media and agents for pharmaceutical active
25 substances is well known in the art. Except insofar as any conventional media or agent is incompatible with the active ingredient, use thereof in the therapeutic compositions is contemplated. Supplementary active ingredients can also be incorporated into the compositions.

30 It is especially advantageous to formulate parenteral compositions in dosage unit form for ease of administration and uniformity of dosage. Dosage unit form as used herein refers to

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physically discrete units suited as unitary dosages for the mammalian subjects to be treated; each unit containing a predetermined quantity of active material calculated to produce the desired therapeutic effect in association with the required pharmaceutical carrier. The specification for the novel dosage unit forms of the invention are dictated by
5 and directly dependent on (a) the unique characteristics of the active material and the particular therapeutic effect to be achieved, and (b) the limitations inherent in the art of compounding such an active material for the treatment of disease in living subjects having a diseased condition in which bodily health is impaired.

10 The principal active ingredient is compounded for convenient and effective administration in effective amounts with a suitable pharmaceutically acceptable carrier in dosage unit form as hereinbefore disclosed. A unit dosage form can, for example, contain the principal active compound in amounts ranging from 0.5 μ g to about 2000 mg. Expressed in proportions, the active compound is generally present in from about 0.5 μ g to about 2000
15 mg/ml of carrier. In the case of compositions containing supplementary active ingredients, the dosages are determined by reference to the usual dose and manner of administration of the said ingredients.

The pharmaceutical composition may also comprise genetic molecules such as a vector
20 capable of transfecting target cells where the vector carries a nucleic acid molecule capable of expressing a sphingosine kinase variant or agent. The vector may, for example, be a viral vector.

Further features of the present invention are more fully described in the following non-
25 limiting examples.

EXAMPLE 1

MATERIALS AND METHODS

5 *Materials* – *D-erythro*-Sphingosine and sphingosine-1-phosphate were purchased from Biomol Research Laboratories Inc. (Plymouth Meeting, PA). ATP and phorbol 12-myristate 13-acetate (PMA) were from Sigma. [$\gamma^{32}\text{P}$] ATP and ^{32}P -phosphoric acid were purchased from Geneworks (Adelaide, South Australia), [choline-methyl- ^{14}C]sphingomyelin from NEN (Boston, MA), TNF α from R&D Systems Inc. (Minneapolis, MN). Interleukin 1 (IL-1) was a gift from Synergyn (Bolder, CO).

Cell Culture and Transfection – Human embryonic kidney cells (HEK293T, ATCC CRL-1573) cells were cultured on Dulbecco's modified Eagle's medium (DMEM; CSL Biosciences, Parkville, Australia) containing 10% fetal calf serum, 2 mM glutamine, 0.2% (w/v) sodium bicarbonate, penicillin (1.2 mg/ml), and gentamycin (1.6 mg/ml). Transfections were performed using the calcium phosphate precipitation method (Graham & van der Eb, 1973). Cells were harvested and lysed by sonication (2 watts for 30 s at 4°C) in lysis buffer containing 50 mM Tris/HCl (pH 7.4), 10% glycerol, 0.05% Triton X-100, 150 mM NaCl, 1 mM dithiothreitol, 2 mM Na₃VO₄ 10 mM NaF, and 1 mM EDTA. Protein concentrations in cell homogenates were determined with either the Coomassie Brilliant Blue (Sigma) or Bichinchoninic acid (Pierce) reagents using BSA as standard.

Enzyme Assays - Sphingosine kinase activity was determined using *D-erythro*-sphingosine and [$\gamma^{32}\text{P}$]ATP as substrates, as described previously (Pitson *et al.*, in press). Neutral sphingomyelinase activity was determined using [choline-methyl- ^{14}C]sphingomyelin as substrate, essentially as previously described (Wiegmann *et al.*, 1994). Briefly, whole cell lysates, prepared as described above, were added to an equal volume of 100 mM Tris/HCl buffer (pH 7.4) containing 0.2% Triton X-100, 10 mM MgCl₂ and [choline-methyl- ^{14}C]sphingomyelin (50,000 cpm/assay) and incubated at 37°C for 60 min. Radioactive phosphorylcholine produced was then extracted with chloroform/methanol (2:1, v/v) and quantified in the aqueous phase by scintillation counting. The measurement of PKC activity *in situ* was performed as described previously (Xia *et al.*, 1996). Briefly, cells were seeded in 24-well plates and maintained in culture medium until 70-80% confluent.

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After the indicated treatments, the cells were washed with DMEM and placed in 60 μ l of buffered salt solution (137 mM NaCl, 5.4 mM KCl, 0.3 mM Na₂HPO₄, 0.4 mM KH₂PO₄, 5.5 mM glucose, and 20 mM HEPES) supplemented with 50 μ g/ml digitonin, 10 mM MgCl₂, 25 mM β -glycerophosphate, and 10 μ M [γ ³²P]ATP (5000 cpm/pmol). A PKC-specific peptide substrate (RKRTLRL) was then added (to 200 μ M) in the presence of 5 mM EGTA and 2.5 mM CaCl₂. After a 10 min incubation at 30°C, the kinase reaction was terminated by the addition of 20 μ l of 25% (w/v) trichloroacetic acid. Aliquots (65 μ l) of the acidified reaction mixtures were spotted on phosphocellulose papers (Whatman P-81) and washed three times with 75 mM phosphoric acid and once with 75 mM sodium phosphate (pH 7.5). The PKC-dependent phosphorylated peptide substrate bound to the filter was quantified by scintillation counting.

Western Blotting - SDS-PAGE was performed on cell lysates according to the method of Laemmli [29] using 12% acrylamide gels. Proteins were blotted to nitrocellulose and the membranes blocked overnight at 4°C in PBS containing 5% skim milk and 0.1% Tris-ton X-100. Sphingosine kinase expression levels were analysed with the M2 anti-FLAG antibody (Sigma). ERK activation in response to agonists was followed in cells serum-starved for 4 h using anti-ERK1/2 (Zymed, San Francisco, CA) and anti-phospho-ERK1/2 (Promega, Madison, WI) antibodies. Immunocomplexes were detected after conjugation to either HRP anti-mouse (Pierce) or anti-rabbit (Selinus/AMRAD, Melbourne, Australia) IgG using an enhanced chemiluminescence kit (Amersham).

Mutagenesis of the SK-1 sequence

The SK-1 cDNA (as described in Pitson *et al.* 2000a) was cloned into pALTER (promega Inc., Madison, WI) site directed mutagenesis vector. Single-stranded DNA was prepared and used as template for oligonucleotide directed mutagenesis as detailed in the manufacturer's protocol. The mutagenic oligonucleotide (5' CTG GAG ACG ATC TGA TGC AC) [≤ 400] was designed to generate the G82D mutant, substitution of the glycine at position 82 to aspartic acid. The mutant was sequenced to verify incorporation of the desired modification.

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Expression of the G82D cDNA

The G82D mutant cDNA was sub-cloned into pcDNA3 (Invitrogen Corp., San Diego CA).

The expression construct was transfected by calcium phosphate precipitation into

5 HEK293T cells.

Transformation assay

For focus formation assay, low passage NIH 3T3 cells were transfected with the V12
10 mutant H-ras, v-SRC (gifts from Dr. Julian Downward²), SphK, G82D mutant SphK
expression vectors or empty vector using Lipofectamine Plus as described above. Two
days later, the transfected cells were split to 6-well plates. After reaching confluence, they
were kept for two weeks in DMEM containing 5% calf serum. The foci were visualized and
scored after staining with 0.5% crystal violet. For soft agar assay, suspensions of 1×10^4
15 cells from the stable transfected pools in a growth medium containing 0.33% agar were
overlaid onto 0.6% agar gel in the absence or presence of DMS at various concentrations.
After 14-days incubation colonies were stained with 0.1 mg/ml MTT and those greater
than 0.1 mm in diameter were scored as positive.

20

EXAMPLE 2**RESULTS**

A mutant of sphingosine kinase that is inactive in its capacity to phosphorylate sphingosine
25 to sphingosine-1-phosphate (S1P), the molecule that mediates the biologically relevant
functions of sphingosine kinase, has been designed and made. This mutant was made by
site directed mutagenesis of a putative ATP binding site (G in position 82 to aspartic acid
'G82D'), thus rendering the sphingosine kinase catalytically inactive

30

G82DSK is well expressed as seen in Western blots (Figure 2) of the FLAG tagged
transfectants and is correctly folded as judged by binding to calmodulin (data not shown).

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The G82DSK by itself has no sphingosine kinase activity and does not suppress endogenous baseline sphingosine kinase activity (Fig. 2), however it totally suppresses the increases in sphingosine kinase activity seen after treatment of cells with activating agents such as TNF, IL-1 and PMA (Fig. 3 & 4). This 'duality' of function is also seen in cells that
5 overexpress sphingosine kinase: the overexpressed baseline levels are not altered, but activation is decreased (Fig. 5). The extent of prevention is likely to depend on the molar ratio of sphingosine kinase:G82DSK.

Furthermore, G82DSK inhibits sphingosine kinase stimulated by the oncogene Ras (Fig 6)
10 and may also suppress *in vitro* and *in vivo* markers of oncogenesis. The inhibitor is specific as it does not depress the activation of another enzyme protein kinase C (fig. 7) or sphingomyelinase (Fig. 8). Furthermore, its function is stable as it inhibits TNF mediated activation at all time point (Fig. 4) and inhibits downstream effects of TNF such as activation of erk (Fig. 9).

15 Thus there has been generated an inhibitor of sphingosine kinase that is quite different from the chemical inhibitor hitherto used widely, N,N-dimethylsphingosine (DMS). DMS totally eliminates wild-type sphingosine kinase function whereas G82DSK only eliminates activation of wild-type sphingosine kinase. In a way, we have a prototype for a new kind
20 of drug.

EXAMPLE 3

Human sphingosine kinase (hSK) has been cloned and found, through sequence analysis to
25 have, similarity in amino acids 16 to 153 to the putative catalytic domain of diacylglycerol kinases (DGKs) (Pitson *et al.*, 2000a). Examples 1 and 2 disclose generation of a catalytically inactive mutant of hSK by site-directed mutagenesis of a single amino acid within this region (see also Pitson *et al.*, 2000b). This mutation, Gly⁸² to Asp (referred to as G82D or hSK^{G82D}) was based on similar mutations of DGKs that also produced inactive
30 mutants (Masai *et al.*, 1993; Topham and Prescott, 1999; Topham *et al.*, 1998). The mechanism whereby the point mutation ablated activity in DGKs and hSK has been

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proposed to be through altering the ATP binding site of the enzymes. This is based on the (loose) similarity in the amino acid sequence motifs for ATP binding sites in protein kinases (Hanks *et al.*, 1988; Benner and Gerlo, 1992) and those found in DGKs and hSK. However, until now, no firm data has been generated to support the involvement of Gly⁸² in ATP binding. This Example describes the generation of another hSK mutant through mutagenesis of Gly⁸² to Ala (G82A or hSK^{G82A}).

Materials and Methods

Materials – D-erythro - Sphingosine was purchased from Biomol Research Laboratories Inc. (Plymouth Meeting, PA), ATP from Sigma, and [$\gamma^{32}\text{P}$]ATP from Geneworks (Adelaide, South Australia).

Cell Culture and Transfection – Human embryonic kidney cells (HEK293T, ATCC CRL-1573) cells were cultured on Dulbecco's modified Eagle's medium (DMEM; CSL Biosciences, Parkville, Australia) containing 10% fetal calf serum, 2 mM glutamine, 0.2% (w/v) sodium bicarbonate, penicillin (1.2 mg/ml), and gentamycin (1.6 mg/ml).

Transfections were performed using the calcium phosphate precipitation method (Graham & van der Eb, 1973). Cells were harvested and lysed by sonication (2 watts for 30 s at 4°C) in lysis buffer containing 50 mM Tris/HCl (pH 7.4), 10% glycerol, 0.05% Triton X-100, 150 mM NaCl, 1 mM dithiothreitol, 2 mM Na₃VO₄, 10 mM NaF, and 1 mM EDTA. Protein concentrations in cell homogenates were determined with the Coomassie Brilliant Blue (Sigma) reagent using BSA as standard.

Enzyme Assays - Sphingosine kinase activity was determined using D-erythro-sphingosine and [$\gamma^{32}\text{P}$]ATP as substrates, as described previously (Pitson *et al.*, 2000a). Kinetic parameters were calculated using a non-linear regression program.

Construction of SK^{G82A} – The SKI cDNA (as described in Pitson *et al.*, 2000a) was cloned into pALTER (Promega Inc., Madison, WI) site directed mutagenesis vector. Single-stranded DNA was prepared and used as a template for oligonucleotide directed mutagenesis as detailed in the manufacturer's protocol. The mutagenic oligonucleotide (5'-

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GTCTGGAGATGCATTGATGCACG-3') was designed to generate the SK^{G82A} mutant, substitution of the glycine at position 82 to alanine. The mutant was sequenced to verify incorporation of the desired modification and sub-cloned into pcDNA3 (Invitrogen Corp., San Diego CA) for expression in HEK293T cells.

5

Western Blotting – SDS-PAGE was performed on cell lysates according to the method of Laemmli (1970) using 12% acrylamide gels. Proteins were blotted to nitrocellulose and the membranes blocked overnight at 4°C in PBS containing 5% skim milk and 0.1% Triton X-100. Sphingosine kinase expression levels were analysed with the M2 anti-FLAG antibody (Sigma). ERK activation in response to agonists was followed in cells serum-starved for 4 h using anti-ERK1/2 (Zymed, San Francisco, CA) and anti-phospho-ERK1/2 (Promega, Madison, WI) antibodies. Immunocomplexes were detected after conjugation to either HRP anti-mouse (Pierce) or anti-rabbit)(Selinus/AMRAD, Melbourne, Australia) IgG using an enhanced chemiluminescence kit (Amersham).

15

EXAMPLE 4

RESULTS

In contrast to the hSK^{G82D} mutant, hSK^{G82A} has catalytic activity, albeit much lower (ca 20 5%) than the wildtype hSK (Figure 12). Analysis of the substrate kinetics of hSK^{G82A} has shown that this mutant has considerably lower affinity for ATP than the wildtype hSK (Figure 13), while the affinity for sphingosine remains unaffected (Figure 14). This kinetic data, summarised in Table 3, indicate that Gly⁸² is involved in ATP binding and suggests that this residue may be part of the ATP-binding site of hSK. This provides evidence that 25 the original Gly⁸² to Asp mutation in hSK ablates catalytic activity of hSK by interruption of ATP binding.

EXAMPLE 5

CONSTRUCTION OF hSK MUTUANTS

Construction of hSK mutants – The SK1 cDNA (as described in Pitson *et al.*, 2000a) was
5 cloned into pALTER (Promega Inc., Madison, WI) site directed mutagenesis vector.
Single-stranded DNA was prepared and used as a template for oligonucleotide directed
mutagenesis as detailed in the manufacturer's protocol. The mutagenic oligonucleotides
used to generate the hSK mutants are listed in Table 1. The mutants were sequenced to
verify incorporation of the desired modification and sub-cloned into pcDNA3 (Invitrogen
10 Corp., San Diego CA) for expression in HEK293T cells.

Those skilled in the art will appreciate that the invention described herein is susceptible to
variations and modifications other than those specifically described. It is to be understood
that the invention includes all such variations and modifications. The invention also
15 includes all of the steps, features, compositions and compounds referred to or indicated in
this specification, individually or collectively, and any and all combinations of any two or
more of said steps or features.

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Table 3 Substrate kinetics of hSK^{WT} and hSK^{G82A}

Kinetic values were obtained by non-linear regression analysis of data presented in Figures 12 and 13. V_{\max} is expressed as a percentage of the maximum velocity calculated for SK^{WT} and standardised for the expression levels of the two recombinant proteins.

	K_m		V_{\max}
	ATP	Sphingosine	
hSK^{WT}	115 μM	12.1 μM	100
hSK^{G82A}	4.2 mM	15.5 μM	32

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Table 4 Mutagenic oligonucleotides used for site-directed mutagenesis of hSK

Mismatches with the hSK^{WT} template are indicated by lowercase letters.

Name	Sequence
G26A	GAACCCGCGGGGCGCCAAGGGCAA (<400>13)
G26D	GCTGAACCCCCGGGGCGACAAGGGCAA (<400>14)
K27A	CGCGGCGGCGCCGGCAAGGCC (<400>15)
K29A	GGCAAGGGCGCCGCCTTGACG (<400>16)
S79A	GTGGTCATGGCCGGCGACGGGCTG (<400>17)
S79D	GTGGTCATGGATGGAGACGGCCTGATGCAC (<400>18)
G80A	TCATGTCTGCAGACGGGCT (<400>19)
G80D	TCATGTCTGACGACGGCCTGATGCAC (<400>20)
G82A	GTCTGGAGATGCATTGATGCACG (<400>21)
G82D	CTGGAGACGATCTGATGCAC (<400>22)
K103A	GCCATCCAGGCCCCCCTGTGT (<400>23)
K103R	GCCATCCAGCGGCCGCTGTGTAGC (<400>24)
G111A	AGCCTCCCTGCAGCCTCTGGCAA (<400>25)
G111D	TCCCAGCAGACTCTGGCAA (<400>26)
G113D	CCCAGCAGGATCCGACAACGCGCT (<400>27)

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CLAIMS:

1. A sphingosine kinase variant comprising a mutation in a region defined by amino acids 16-153 or functionally equivalent region wherein said variant exhibits ablated or reduced catalytic activity relative to wild-type sphingosine kinase or a derivative, homologue, analogue, chemical equivalent or mimetic of said sphingosine kinase variant.
2. The sphingosine kinase variant according to claim 1 wherein said sphingosine kinase is human sphingosine kinase.
3. The sphingosine kinase variant according to claim 1 or 2 wherein said mutation comprises a single or multiple amino acid substitution, addition and/or deletion.
4. The sphingosine kinase variant according to claim 3 wherein said region is defined by amino acids 70-90.
5. The sphingosine kinase variant according to claim 4 wherein said region is defined by amino acids 79-84.
6. The sphingosine kinase variant according to claim 5 wherein said mutation is an amino acid substitution of the glycine amino acid at position 82 to aspartic acid.
7. The sphingosine kinase variant according to any one of claims 3-5 wherein said variant exhibits ablated catalytic activity.
8. The sphingosine kinase variant according to claim 7 wherein said variant comprises one or more of the amino acid substitutions selected from the following list:
 - (i) G82D
 - (ii) G82A
 - (iii) G26D
 - (iv) S79D

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- (v) G80D
- (vi) K103A
- (vii) G111D
- (viii) G113D
- (ix) G26A
- (x) K27A
- (xi) K29A
- (xii) S79A
- (xiii) G80A
- (xiv) K103R
- (xv) G111A

9. A sphingosine kinase variant comprising a mutation in an ATP binding site region or functionally equivalent region wherein said variant exhibits ablated or reduced catalytic activity relative to wild-type sphingosine kinase or a derivative, homologue, analogue, chemical equivalent or mimetic of said sphingosine kinase variant.
10. The sphingosine kinase variant according to claim 9 wherein said sphingosine kinase is human sphingosine kinase.
11. The sphingosine kinase variant according to claim 9 or 10 wherein said mutation comprises a single or multiple amino acid substitution, addition and/or deletion.
12. The sphingosine kinase variant according to claim 11 wherein said region is defined by amino acids 70-90.
13. The sphingosine kinase variant according to claim 12 wherein said region is defined by amino acids 79-84.

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14. The sphingosine kinase variant according to claim 13 wherein said mutation is an amino acid substitution of the glycine amino acid at position 82 to aspartic acid.
15. The sphingosine kinase variant according to any one of claims 11-13 wherein said variant exhibits ablated catalytic activity.
16. The sphingosine kinase variant according to claim 15 wherein said variant comprises one or more of the amino acid substitutions selected from the following list:
 - (i) G82D
 - (ii) G82A
 - (iii) G26D
 - (iv) S79D
 - (v) G80D
 - (vi) K103A
 - (vii) G111D
 - (viii) G113D
 - (ix) G26A
 - (x) K27A
 - (xi) K29A
 - (xii) S79A
 - (xiii) G80A
 - (xiv) K103R
 - (xv) G111A
17. An isolated nucleic acid molecule selected from the list consisting of:
 - (i) An isolated nucleic acid molecule or derivative or equivalent thereof comprising a nucleotide sequence encoding or complementary to a sequence encoding a sphingosine kinase variant or derivative, homologue, analogue, chemical equivalent or mimetic of said variant which variant

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comprises a mutation in a region defined by amino acid 16-153 or functionally equivalent region wherein said variant exhibits ablated or reduced catalytic activity relative to wild-type sphingosine kinase.

- (ii) An isolated nucleic acid molecule or derivative or equivalent thereof comprising a nucleotide sequence encoding or complementary to a sequence encoding a human sphingosine kinase variant or derivative, homologue, analogue, chemical equivalent or mimetic of said variant which variant comprises a mutation in a region defined by amino acid 16-153 or functionally equivalent region wherein said variant exhibits ablated or reduced catalytic activity relative to wild-type human sphingosine kinase.
- (iii) An isolated nucleic acid molecule or derivative or equivalent thereof comprising a nucleotide sequence encoding or complementary to a sequence encoding a human sphingosine kinase variant or derivative, homologue, analogue, chemical equivalent or mimetic of said variant, which variant comprises an amino acid sequence with a single or multiple multiple amino acid substitution, addition and/or deletion in a region defined by amino acid 16-153 or functionally equivalent region wherein said variant exhibits ablated or reduced catalytic activity relative to wild-type sphingosine kinase.
- (iv) An isolated nucleic acid molecule or derivative or equivalent thereof comprising a nucleotide sequence encoding or complementary to a sequence encoding a human sphingosine kinase variant or derivative, homologue, analogue, chemical equivalent or mimetic of said variant, which variant comprises an amino acid sequence with a single or multiple amino acid substitution, addition and/or deletion in a region defined by amino acid 70-90 or functionally equivalent region wherein said variant exhibits ablated or reduced catalytic activity relative to wild-type sphingosine kinase.

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- (v) An isolated nucleic acid molecule or derivative or equivalent thereof comprising a nucleotide sequence encoding or complementary to a sequence encoding a human sphingosine kinase variant or derivative, homologue, analogue, chemical equivalent or mimetic of said variant, which variant comprises an amino acid sequence with a single or multiple multiple amino acid substitution, addition and/or deletion in a region defined by amino acid 79-84 or functionally equivalent region wherein said variant exhibits ablated or reduced catalytic activity relative to wild-type sphingosine kinase.
- (vi) An isolated nucleic acid molecule or derivative or equivalent thereof comprising a nucleotide sequence encoding or complementary to a sequence encoding a sphingosine kinase variant or a derivative, homologue, analogue, chemical equivalent or mimetic of said variant comprising one or more of the amino acid substitutions selected from the following list:
- (a) G82D
 - (b) G82A
 - (c) G26D
 - (d) S79D
 - (e) G80D
 - (f) K103A
 - (g) G111D
 - (h) G113D
 - (i) G26A
 - (j) K27A
 - (k) K29A
 - (l) S79A
 - (m) G80A
 - (n) K103R
 - (o) G111A

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- (vii) An isolated nucleic acid molecule or derivative or analogue thereof comprising a nucleotide sequence encoding or complementary to a sequence encoding a sphingosine kinase variant or derivative, homologue, analogue, chemical equivalent or mimetic of said variant which variant comprises a mutation in an ATP binding site region or functionally equivalent region wherein said variant exhibits ablated or reduced catalytic activity relative to wild-type sphingosine kinase.
-
- 18. A method for detecting an agent capable of modulating the interaction of FOSK with sphingosine kinase or its functional equivalent or derivative thereof said method comprising contacting a cell or extract thereof containing said sphingosine kinase and FOSK or its functional equivalent or derivative with a putative agent and detecting an altered expression phenotype associated with said interaction.
 - 19. A method for detecting an agent capable of binding or otherwise associating with the sphingosine kinase region defined by amino acid 16-153 or functional equivalent or derivative thereof said method comprising contacting a cell containing said amino acid region or functional equivalent or derivative thereof with a putative agent and detecting an altered expression phenotype associated with modulation of the function of sphingosine kinase or its functional equivalent or derivative.
 - 20. The method according to claim 19 wherein said amino acid region is defined by amino acid 70-90.
 - 21. The method according to claim 20 wherein said amino acid region is defined by amino acid 79-84.
 - 22. A method for analysing, designing and/or modifying an agent capable of interacting with the sphingosine kinase region defined by amino acid 16-153 or derivative thereof and modulating at least one functional activity associated with

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said sphingosine kinase said method comprising contacting said sphingosine kinase or derivative thereof with a putative agent and assessing the degree of interactive complementarity of said agent with said binding site.

23. The method according to claim 22 wherein said amino acid region is defined by amino acid 70-90.
24. The method according to claim 23 wherein said amino acid region is defined by amino acid 79-84.
25. The agent identified in accordance with the method of any one of claims 18-24.
26. A method of modulating cellular functional activity in a mammal said method comprising administering to said mammal an effective amount of a sphingosine kinase variant according to any one of claims 1-17 or agent according to claim 25 for a time and under conditions sufficient to inhibit, reduce or otherwise down-regulate at least one functional activity of wild-type sphingosine kinase.
27. The method according to claim 26 wherein said activity is down-regulation of wild-type sphingosine kinase baseline activity and/or prevention of wild-type sphingosine kinase activation.
28. The treatment and/or prophylaxis of a condition in a mammal, which condition is characterised by aberrant, unwanted or otherwise inappropriate cellular activity, said method comprising administering to said mammal an effective amount of a sphingosine kinase variant according to any one of claims 1-17 or agent according to claim 25 for a time and under conditions sufficient to inhibit, reduce or otherwise down-regulate at least one functional activity of wild-type sphingosine kinase wherein said down-regulation results in modulation of cellular functional activity.

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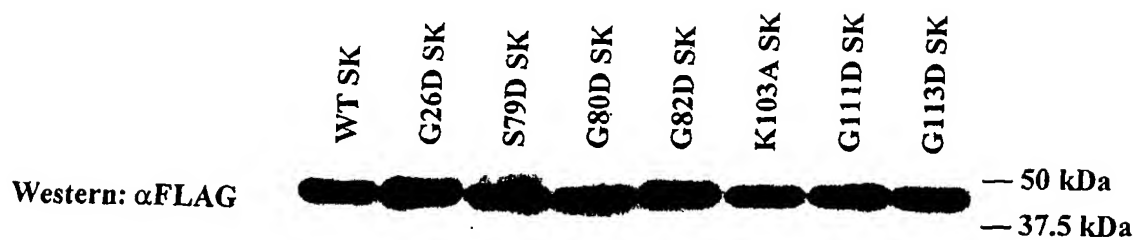
29. The method according to claim 28 wherein said activity is down-regulation of wild-type sphingosine kinase baseline activity and/or prevention of wild-type sphingosine kinase activation.
30. The use of a sphingosine kinase variant according to any one of claims 1-17 or agent according to claim 25 in the manufacture of a medicament for the modulation of cellular functional activity.
31. A sphingosine kinase variant according to any one of claims 1-17 or agent according to claim 25 for use in modulating cellular functional activity.
32. A pharmaceutical composition comprising a sphingosine kinase variant according to any one of claims 1-17 or agent according to claim 25 together with one or more pharmaceutically acceptable carriers and/or diluents.

Figure 1

<400>1	Drosophila DGK2	809	PVIVFIRKKGKNGOGH-KLIGKFECHLIPNPROV-FDLTQ-GGPKMGIDM-ERKAPNL---	RYLAC	871
<400>2	Human DGK ₅	295	PLIFVIRKKGKNGOGH-KIIQISFLWYNBPRO-FDLISQ-GGPKKALEM-YRKVHNL---	RILAC	357
<400>3	Human DGK ₆	219	PLIILASRKTNNNGE-GLIGFEIILNPNQV-FDVTK-TPPIKALQL-CTLLPYSA-RVLVC		282
<400>4	Human SK1	16	RVILVIRKKGKALQLFRSHVQVLPAEAEISFTLMTERRNHAREL-VRSEELGRDALVMS	LM	84
<400>5	Mouse SK1	22	RVILVIRKKGKALQLFQSRVQPELEAEITFKLITERKNHAREL-VCAEELGHWDALAVMS	LM	90
<400>6	Human SK2	146	RLILVIRKKGKALQFQSRVQPELEAEITFKLITERKNHAREL-VOGLSISEWDGIYVS	LL	214
<400>7	Mouse SK2	147	RLILVIRKKGKALQFQSRVQPELEAEITFKLITERKNHAREL-VOGLSISEWDGIYVS	LL	215
<400>8	Yeast LCB4	228	SLILVIRKKGKAKNLELTAKPIVESGCKIETAYTKYARHAIID-AKDLDISKYDTIACAS	LP	296
<400>9	Yeast LCB5	270	SILVIRKKGKAKNLELTAKPIVESGCKIETAYTKYARHAIID-AKDLDISKYDTIACAS	LP	338
<400>10	S. pombe SK	107	RFLVIRKKGKAKHWESEAEVFFSSAHSICEVVLTRKDHAKSI-AKNLDVGSYDGIISV	LF	175
<400>11	C. elegans SK	85	NLILVIRKKGKAKHWESEAEVFFSSAHSICEVVLTRKDHAKSI-AKNLDVGSYDGIISV	LF	154
<400>12	Arabidopsis SK	279	RLILVIRKKGKAKHWESEAEVFFSSAHSICEVVLTRKDHAKSI-AKNLDVGSYDGIISV	IL	347

Drosophila DGK2	872	GWILSVLDQIQPPLQ-PALAVGVLLD	PARAGWGGGYTD---	EPIGKIILREIG-MSQCVLMDR	935			
Human DGK ₅	358	GWILSTLDQRLKP----	PEIVAILL	PARAGWGGGYTD---	EPVSKILSHVE-EGNVQQLDR	419		
Human DGK ₆	283	GWILDAVDNMKIKGQEKYILQVAVLL	D	SNTGWGTGYAG---	EIPVQVLANVMEADGIKIDR	350		
Human SK1	85	HEVINGLMERPDWETAI-OKLCSL	AS	PAAS	NHYAGYEQVTNEDLLTNC	CTLLCRRLSPNNLL	153	
Mouse SK1	91	HEVINGLMERPDWETAI-OKLCSL	AS	PAAS	NHYAGYEQVTNEDLLTNC	CTLLCRRLSPNNLL	159	
Human SK2	215	HEVINGLIDRPDWEAV-KMVGIL	CS	PAAGAVN	HOGGFEFALGLDILLNCS	LLCRGGHPDLDL	283	
Mouse SK2	216	YEINGLIDRPDWEAV-KMVGIL	CS	PAAGAVN	HOGGFEFALGLDILLNCS	LLCRGGSHPLDL	284	
Yeast LCB4	297	YEINGLYRRPDRVDAENKLAVTOL	CS	PAAGAVSHHGGFEQVGV	DILLNCSLLCRGGSHPLDL		360	
Yeast LCB5	339	HEINGLYRRPDRVDAENKLAVTOL	CS	PAAGAVSHHGGFEQVGV	DILLNCSLLCRGGSHPLDL		403	
S. pombe SK	176	HEINGLGERDDYLEAF-KLVCMIS	SS	PAEFSYN-----	CHWTNPNPSYALCLVKS	IEIRIDLMCCS	238	
C. elegans SK	154	FEALINGLGERDDYLEAF-KLVCMIS	SS	PAEFSYN-----	ATGQLKPALEIILKGRPTS	FDLMTFE	226	
Arabidopsis SK	348	VEVINGLIERADWRNAL-KLIGMV	PA	TL	GMIKSL	LDIVGLRCCAN SATISILIRG	HKRSVDVATIAQ	515

A. Expression of hSK point mutants



B. Enzyme activity of hSK point mutants

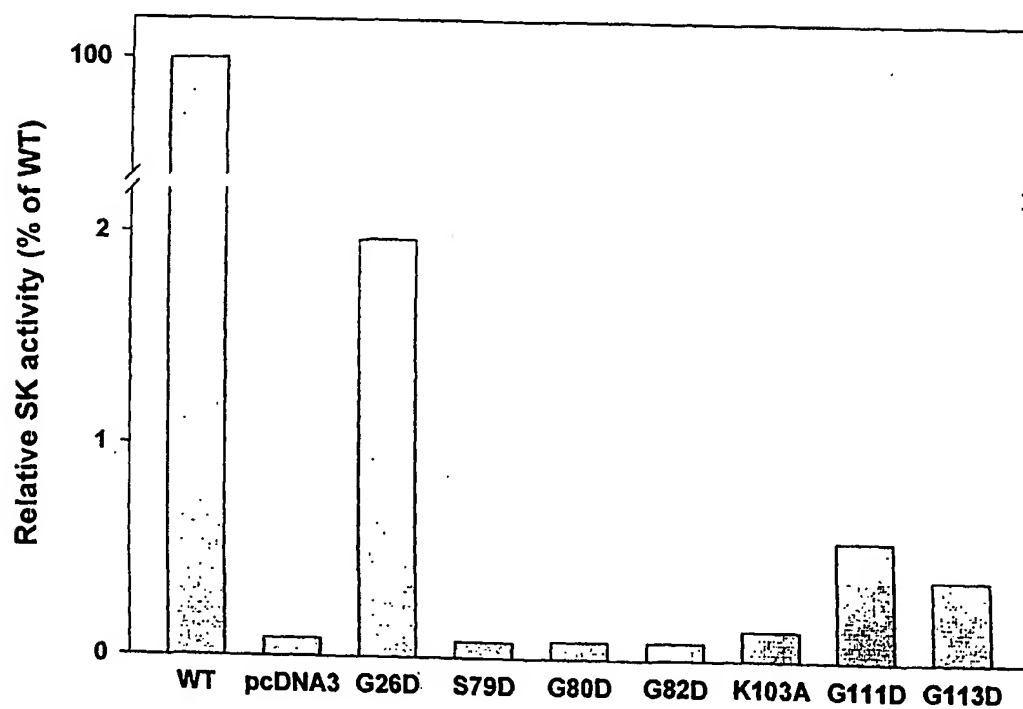


FIGURE 2

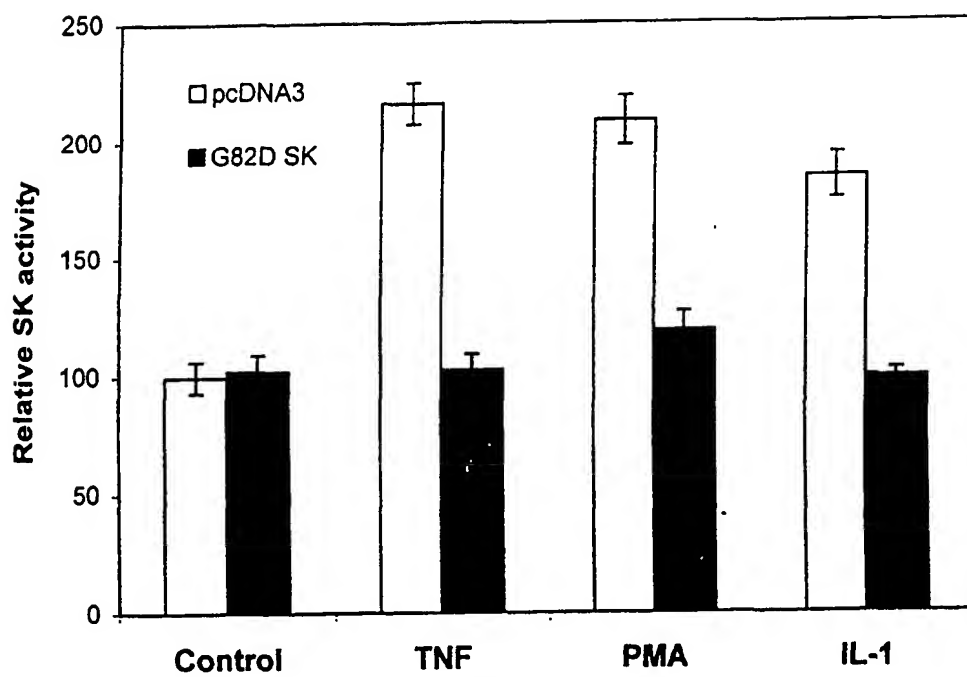
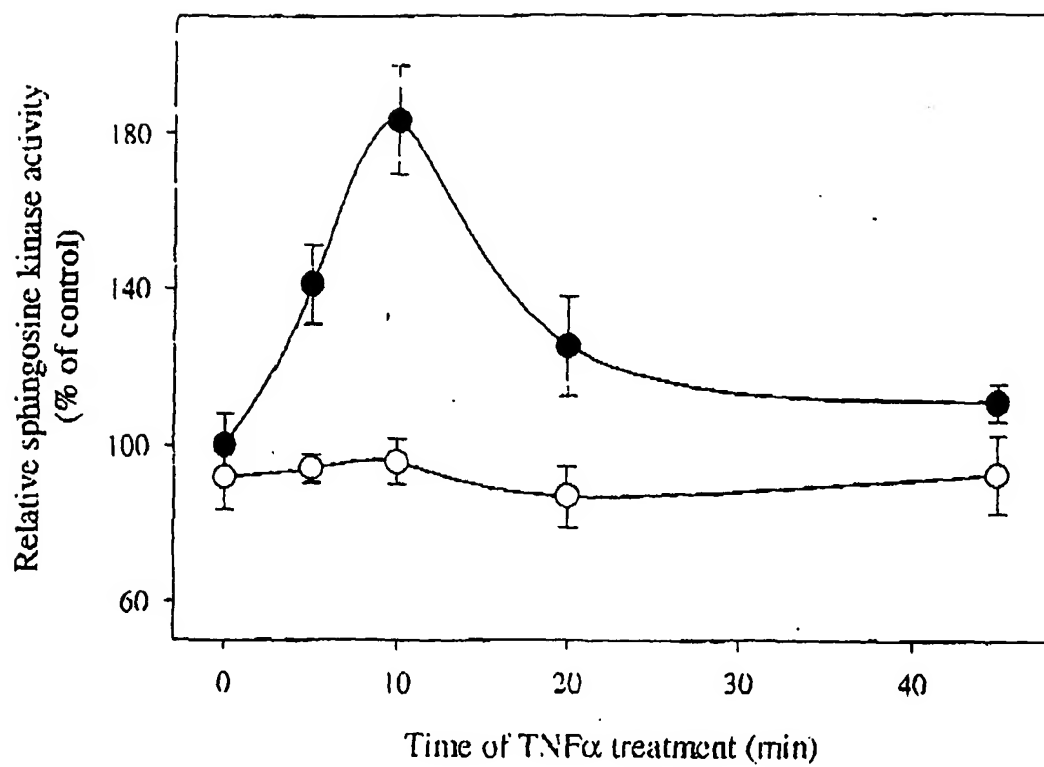


FIGURE 3

FIGURE 4



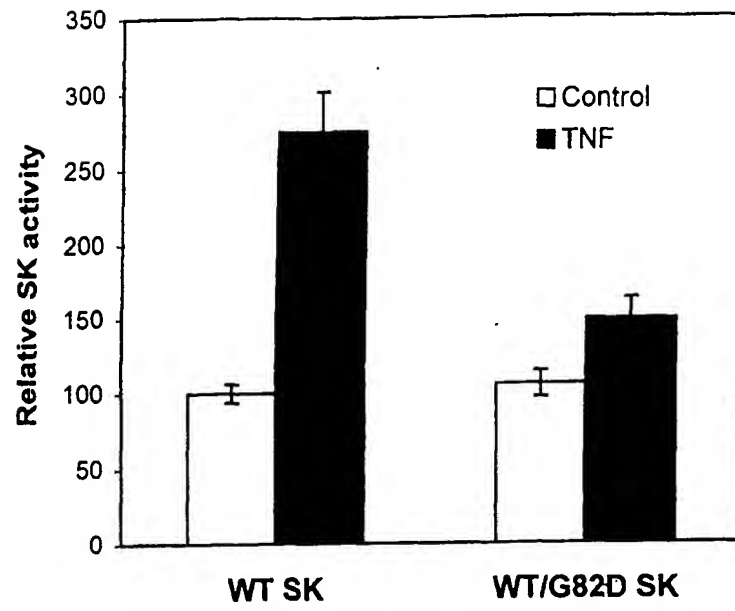


FIGURE 5

Effect of Ras on SK activation

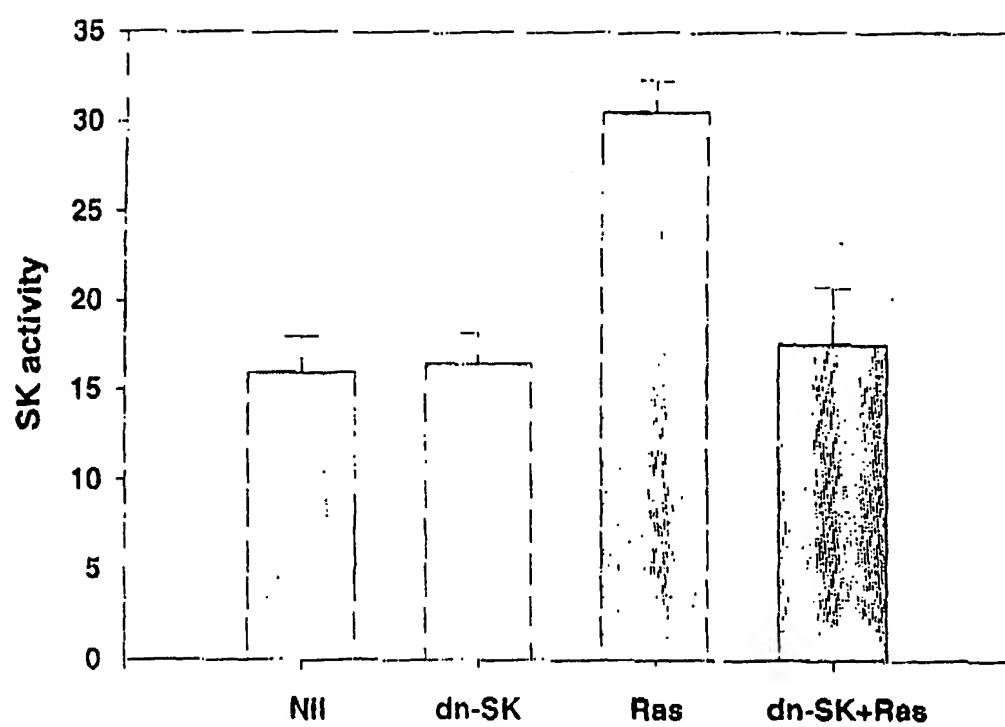


FIGURE 6

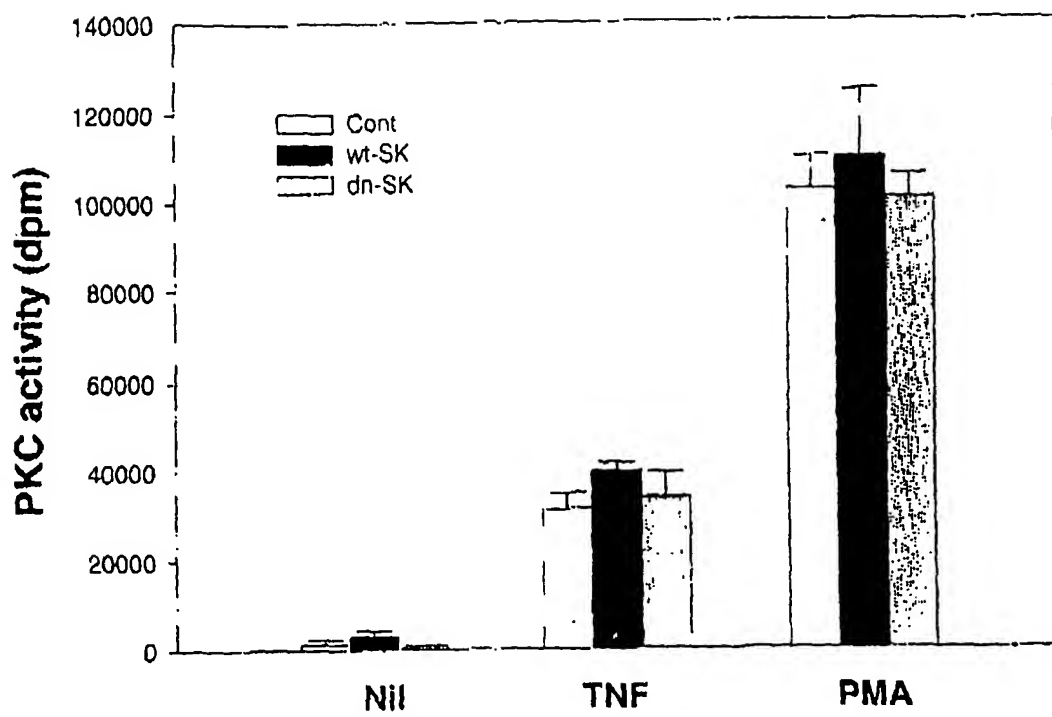


FIGURE 7

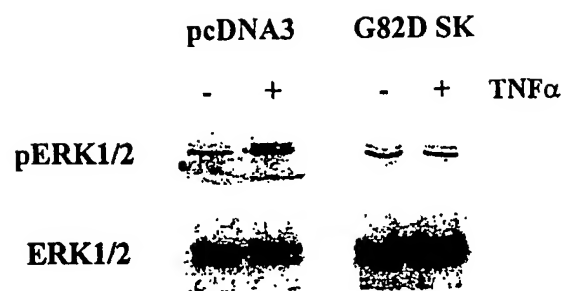
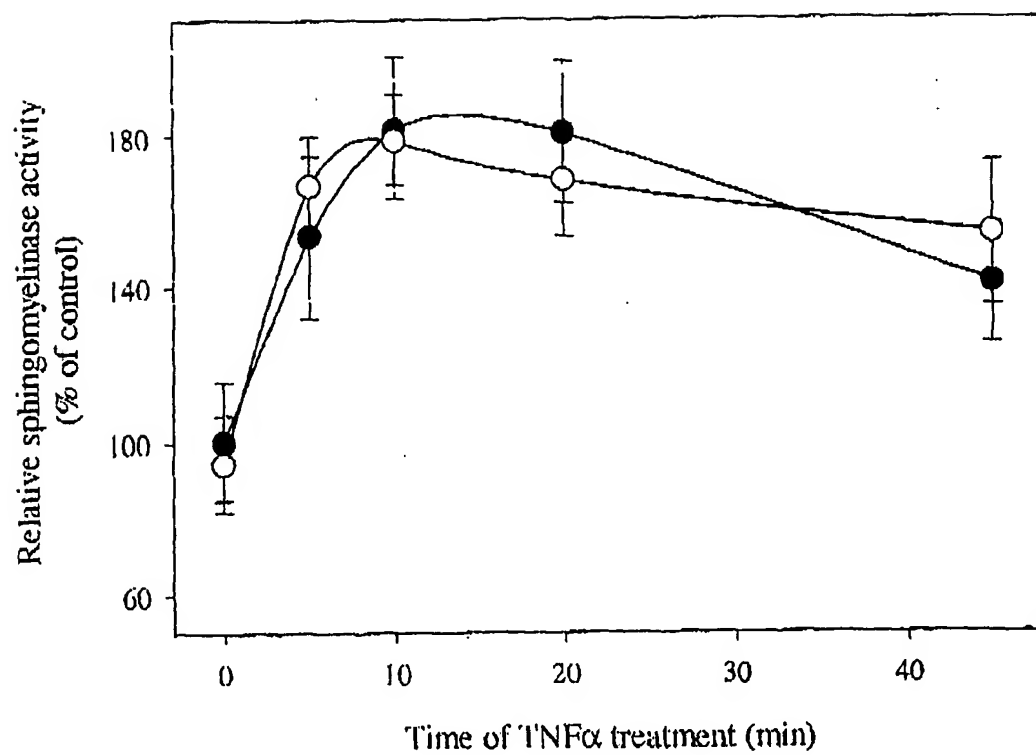
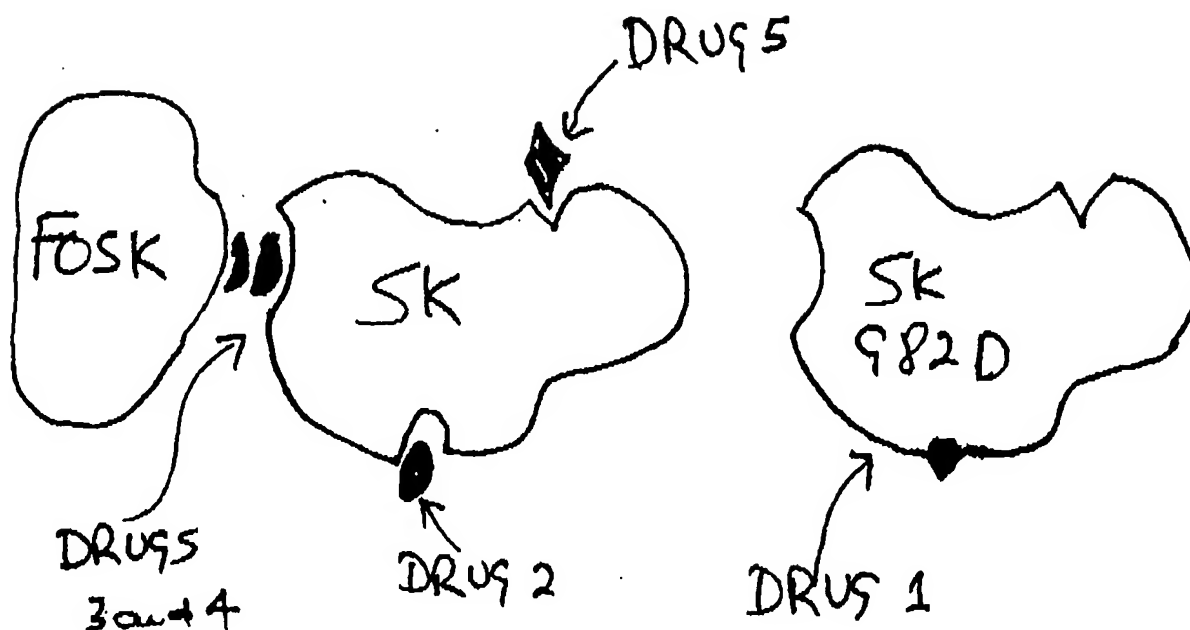


FIGURE 8

FIGURE 9



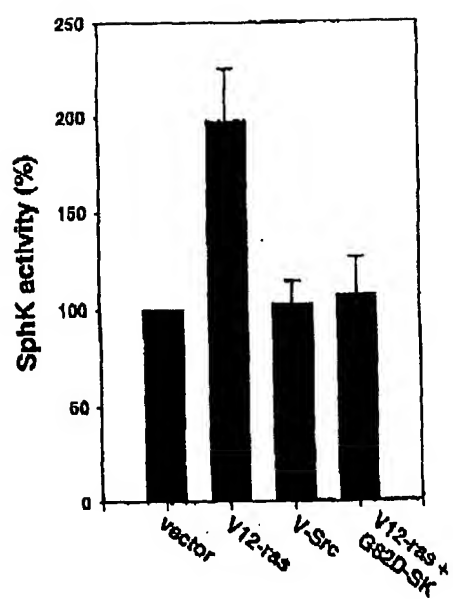


SK = sphingosine kinase
 FOSK = friend of sphingosine kinase

- Drug 1** Is G82D SK (or other equivalent mutants of SK)
- Drug 2** Is a molecule that binds to an area of SK near G82D and renders the molecule inactive. This type of drug will not only render SK inactive, but will also cause the molecule to become a competitor for binding to FOSK, hence also a dominant negative.
- Drugs 3 & 4** Are molecules binding either to SK or FOSK preventing their interaction. These drugs will only inhibit SK activation, not its baseline function. Screening for this type of drug will need to be specialised as the drug will not result in a loss of activity of SK.
- Drug 5** Is a molecule that binds SK and inhibits its function but the molecule will not act as a dominant negative.

FIGURE 10

A.



B.

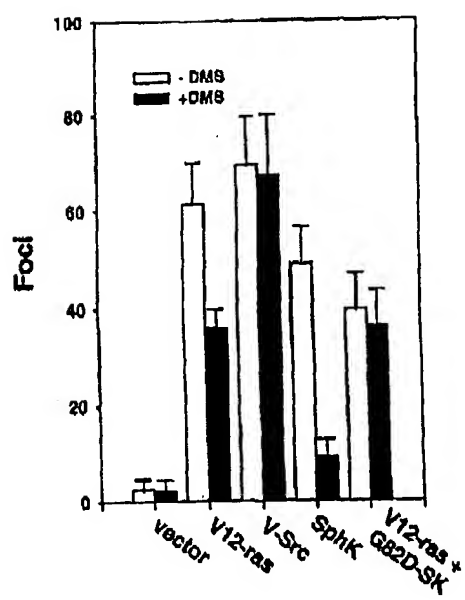


FIGURE 11

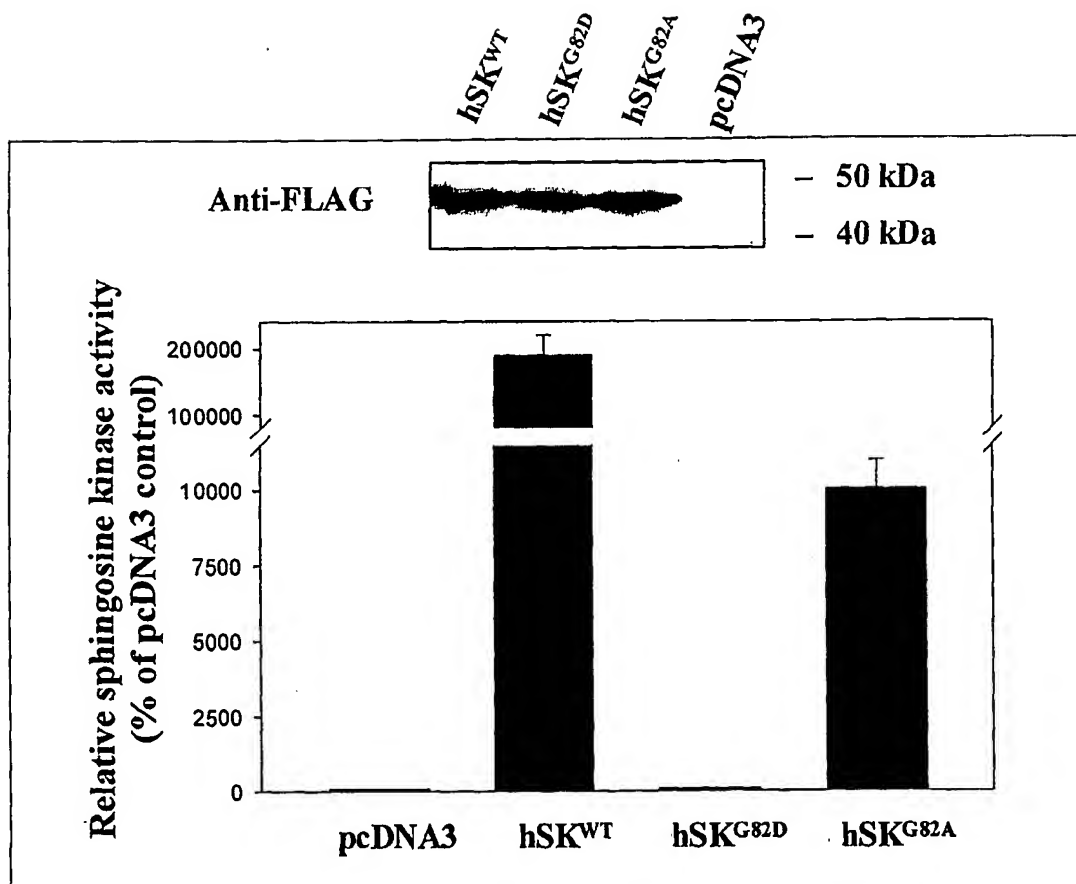


FIGURE 12

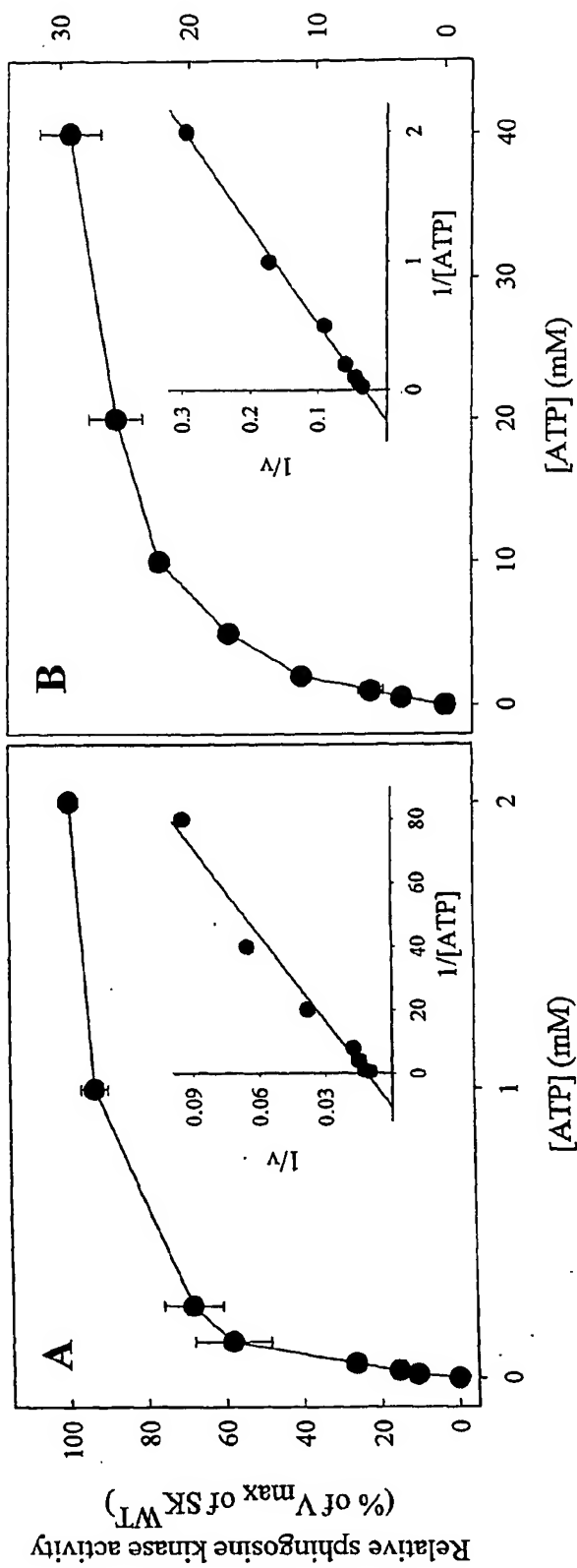


FIGURE 13

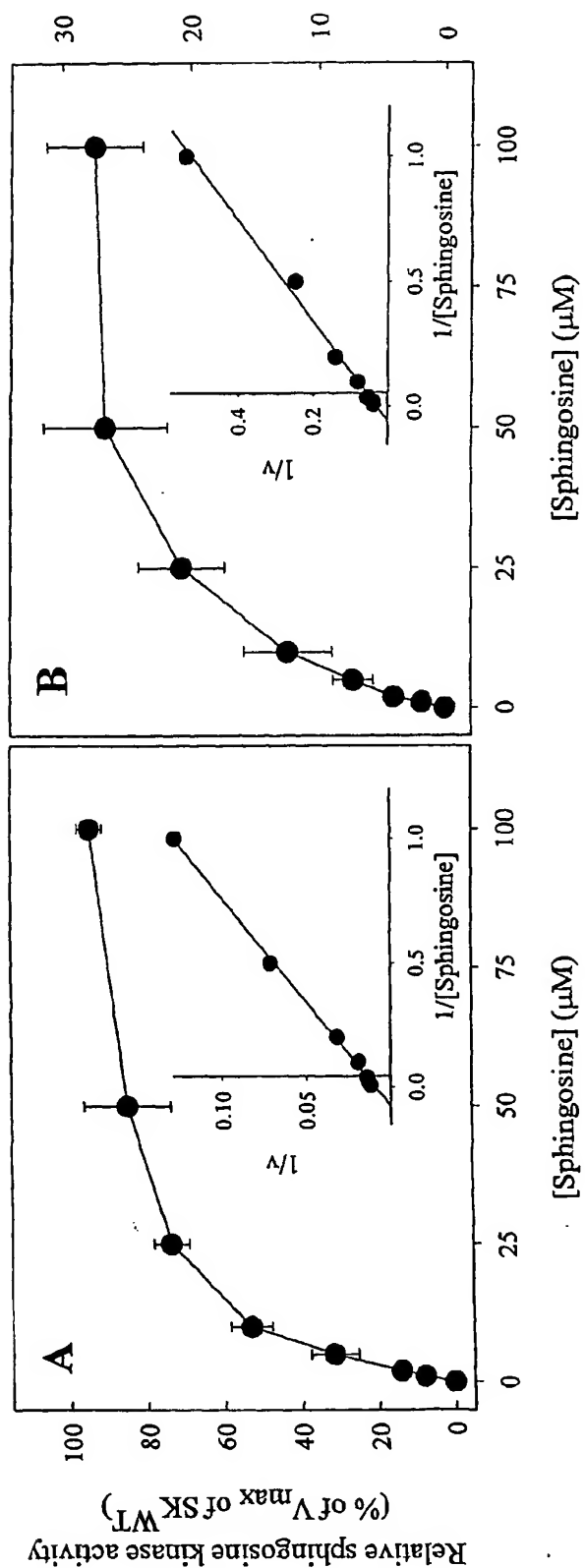


FIGURE 14

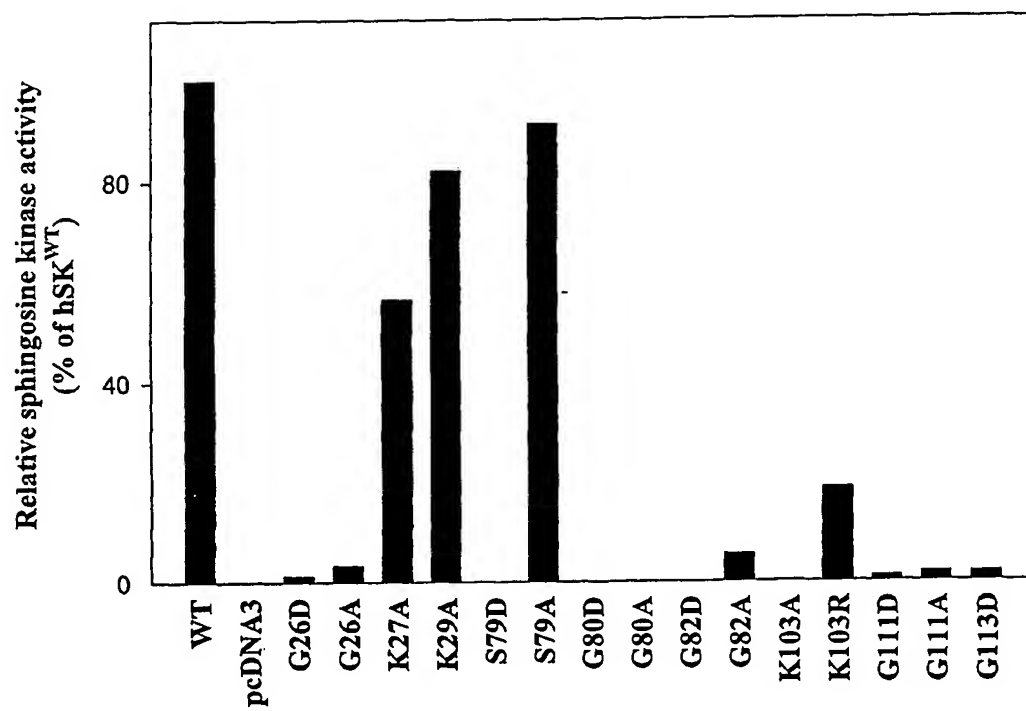


FIGURE 15

- 1 -

SEQUENCE LISTING

<110> Medvet Science Pty Ltd

<120> NOVEL THERAPEUTIC MOLECULAR VARIANTS AND USES THEREOF

<130> 2427140/TDO

<140> International Patent Application

<141> 2001-06-20

<160> 27

<170> PatentIn Ver. 2.0

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<211> 127

<212> PRT

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Lys Leu Leu Gly Lys Phe Gln His Leu Leu Asn Pro Arg Gln Val Phe

20 25 30

Asp Leu Thr Gln Gly Gly Pro Lys Met Gly Leu Asp Met Phe Arg Lys

35 40 45

Ala Pro Asn Leu Arg Val Leu Ala Cys Gly Gly Asp Gly Thr Val Gly

50 55 60

- 2 -

Trp Val Leu Ser Val Leu Asp Gln Ile Gln Pro Pro Leu Gln Pro Ala
 65 70 75 80

Pro Ala Val Gly Val Leu Pro Leu Gly Thr Gly Asn Asp Leu Ala Arg
 85 90 95

Ala Leu Gly Trp Gly Gly Gly Tyr Thr Asp Glu Pro Ile Gly Lys Ile
 100 105 110

Leu Arg Glu Ile Gly Met Ser Gln Cys Val Leu Met Asp Arg Trp
 115 120 125

<210> 2

<211> 125

<212> PRT

<213> mammalian

<400> 2

Pro Leu Leu Val Phe Val Asn Pro Lys Ser Gly Gly Asn Gln Gly Ala
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Lys Ile Ile Gln Ser Phe Leu Trp Tyr Leu Asn Pro Arg Gln Val Phe
 20 25 30

Asp Leu Ser Gln Gly Gly Pro Lys Glu Ala Leu Glu Met Tyr Arg Lys
 35 40 45

Val His Asn Leu Arg Ile Leu Ala Cys Gly Gly Asp Gly Thr Val Gly
 50 55 60

- 3 -

Trp Ile Leu Ser Thr Leu Asp Gln Leu Arg Leu Lys Pro Pro Pro Pro
65 70 75 80

Val Ala Ile Leu Pro Leu Gly Thr Gly Asn Asp Leu Ala Arg Thr Leu
85 90 95

Asn Trp Gly Gly Gly Tyr Thr Asp Glu Pro Val Ser Lys Ile Leu Ser

100 105 110

His Val Glu Glu Gly Asn Val Val Gln Leu Asp Arg Trp
115 120 125

<210> 3

<211> 132

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1 5 10 15

Gly Leu Leu Gly Glu Phe Arg Ile Leu Leu Asn Pro Val Gln Val Phe
20 25 30

Asp Val Thr Lys Thr Pro Pro Ile Lys Ala Leu Gln Leu Cys Thr Leu
35 40 45

Leu Pro Tyr Tyr Ser Ala Arg Val Leu Val Cys Gly Gly Asp Gly Thr
50 55 60

- 4 -

Val Gly Trp Val Leu Asp Ala Val Asp Asp Met Lys Ile Lys Gly Gln
 65 70 75 80

Glu Lys Tyr Ile Pro Gln Val Ala Val Leu Pro Leu Gly Thr Gly Asn
 85 90 95

Asp Leu Ser Asn Thr Leu Gly Trp Gly Thr Gly Tyr Ala Gly Glu Ile
 100 105 110

Pro Val Ala Gln Val Leu Arg Asn Val Met Glu Ala Asp Gly Ile Lys
 115 120 125

Leu Asp Arg Trp
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<210> 4

<211> 138

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<213> mammalian

<400> 4

Arg Val Leu Val Leu Leu Asn Pro Arg Gly Gly Lys Gly Lys Ala Leu
 1 5 10 15

Gln Leu Phe Arg Ser His Val Gln Pro Leu Leu Ala Glu Ala Glu Ile
 20 25 30

Ser Phe Thr Leu Met Leu Thr Glu Arg Arg Asn His Ala Arg Glu Leu
 35 40 45

- 5 -

Val Arg Ser Glu Glu Leu Gly Arg Trp Asp Ala Leu Val Val Met Ser

50

55

60

Gly Asp Gly Leu Met His Glu Val Val Asn Gly Leu Met Glu Arg Pro

65

70

75

80

Asp Trp Glu Thr Ala Ile Gln Lys Pro Leu Cys Ser Leu Pro Ala Gly

85

90

95

Ser Gly Asn Ala Leu Ala Ala Ser Leu Asn His Tyr Ala Gly Tyr Glu

100

105

110

Gln Val Thr Asn Glu Asp Leu Leu Thr Asn Cys Thr Leu Leu Leu Cys

115

120

125

Arg Arg Leu Leu Ser Pro Met Asn Leu Leu

130

135

<210> 5

<211> 138

<212> PRT

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<400> 5

Arg Val Leu Val Leu Leu Asn Pro Gln Gly Gly Lys Gly Lys Ala Leu

1

5

10

15

Gln Leu Phe Gln Ser Arg Val Gln Pro Phe Leu Glu Glu Ala Glu Ile

20

25

30

- 6 -

Thr Phe Lys Leu Ile Leu Thr Glu Arg Lys Asn His Ala Arg Glu Leu

35

40

45

Val Cys Ala Glu Glu Leu Gly His Trp Asp Ala Leu Ala Val Met Ser

50

55

60

Gly Asp Gly Leu Met His Glu Val Val Asn Gly Leu Met Glu Arg Pro

65

70

75

80

Asp Trp Glu Thr Ala Ile Gln Lys Pro Leu Cys Ser Leu Pro Gly Gly

85

90

95

Ser Gly Asn Ala Leu Ala Ala Ser Val Asn His Tyr Ala Gly Tyr Glu

100

105

110

Gln Val Thr Asn Glu Asp Leu Leu Ile Asn Cys Thr Leu Leu Leu Cys

115

120

125

Arg Arg Arg Leu Ser Pro Met Asn Leu Leu

130

135

<210> 6

<211> 138

<212> PRT

<213> mammalian

<400> 6

Arg Leu Leu Leu Leu Val Asn Pro Phe Gly Gly Arg Gly Leu Ala Trp

1

5

10

15

- 7 -

Gln Trp Cys Lys Asn His Val Leu Pro Met Ile Ser Glu Ala Gly Leu
 20 25 30

Ser Phe Asn Leu Ile Gln Thr Glu Arg Gln Asn His Ala Arg Glu Leu
 35 40 45

Val Gln Gly Leu Ser Leu Ser Glu Trp Asp Gly Ile Val Thr Val Ser
 50 55 60

Gly Asp Gly Leu Leu His Glu Val Leu Asn Gly Leu Leu Asp Arg Pro
 65 70 75 80

Asp Trp Glu Glu Ala Val Lys Met Pro Val Gly Ile Leu Pro Cys Gly
 85 90 95

Ser Gly Asn Ala Leu Ala Gly Ala Val Asn Gln His Gly Gly Phe Glu
 100 105 110

Pro Ala Leu Gly Leu Asp Leu Leu Leu Asn Cys Ser Leu Leu Leu Cys
 115 120 125

Arg Gly Gly Gly His Pro Leu Asp Leu Leu
 130 135

<210> 7

<211> 138

<212> PRT

<213> mammalian

- 8 -

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Arg Leu Leu Ile Leu Val Asn Pro Phe Gly Gly Arg Gly Leu Ala Trp
1 5 10 15

Gln Arg Cys Met Asp His Val Val Pro Met Ile Ser Glu Ala Gly Leu
20 25 30

Ser Phe Asn Leu Ile Gln Thr Glu Arg Gln Asn His Ala Arg Glu Leu
35 40 45

Val Gln Gly Leu Ser Leu Ser Glu Trp Glu Gly Ile Val Thr Val Ser
50 55 60

Gly Asp Gly Leu Leu Tyr Glu Val Leu Asn Gly Leu Leu Asp Arg Pro
65 70 75 80

Asp Trp Glu Asp Ala Val Arg Met Pro Ile Gly Val Leu Pro Cys Gly
85 90 95

Ser Gly Asn Ala Leu Ala Gly Ala Val Ser His His Gly Gly Phe Glu
100 105 110

Gln Val Val Gly Val Asp Leu Leu Leu Asn Cys Ser Leu Leu Leu Cys
115 120 125

Arg Gly Gly Ser His Pro Leu Asp Leu Leu
130 135

- 9 -

<210> 8

<211> 133

<212> PRT

<213> Yeast

<400> 8

Ser Ile Leu Val Ile Ile Asn Pro His Gly Gly Lys Gly Thr Ala Lys

1 5 10 15

Asn Leu Phe Leu Thr Lys Ala Arg Pro Ile Leu Val Glu Ser Gly Cys

20 25 30

Lys Ile Glu Ile Ala Tyr Thr Lys Tyr Ala Arg His Ala Ile Asp Ile

35 40 45

Ala Lys Asp Leu Asp Ile Ser Lys Tyr Asp Thr Ile Ala Cys Ala Ser

50 55 60

Gly Asp Gly Ile Pro Tyr Glu Val Ile Asn Gly Leu Tyr Arg Arg Pro

65 70 75 80

Asp Arg Val Asp Ala Phe Asn Lys Leu Ala Val Thr Gln Leu Pro Cys

85 90 95

Gly Ser Gly Asn Ala Met Ser Ile Ser Cys His Trp Thr Asn Asn Pro

100 105 110

Ser Tyr Ala Ala Leu Cys Leu Val Lys Ser Ile Glu Thr Arg Ile Asp

115 120 125

Leu Met Cys Cys Ser

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<211> 133

<212> PRT

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<400> 9

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Lys Leu Phe Met Thr Lys Ala Lys Pro Leu Leu Leu Ala Ser Arg Cys

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Ser Ile Glu Val Val Tyr Thr Lys Tyr Pro Gly His Ala Ile Glu Ile

35 40 45

Ala Arg Glu Met Asp Ile Asp Lys Tyr Asp Thr Ile Ala Cys Ala Ser

50 55 60

Gly Asp Gly Ile Pro His Glu Val Ile Asn Gly Leu Tyr Gln Arg Pro

65 70 75 80

Asp His Val Lys Ala Phe Asn Asn Ile Ala Ile Thr Glu Ile Pro Cys

85 90 95

Gly Ser Gly Asn Ala Met Ser Val Ser Cys His Trp Thr Asn Asn Pro

100 105 110

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115 120 125

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Leu Met Cys Cys Ser

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<213> S. pombe

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His Ile Trp Glu Ser Glu Ala Glu Pro Val Phe Ser Ser Ala His Ser

20 25 30

Ile Cys Glu Val Val Leu Thr Arg Arg Lys Asp His Ala Lys Ser Ile

35 40 45

Ala Lys Asn Leu Asp Val Gly Ser Tyr Asp Gly Ile Leu Ser Val Gly

50 55 60

Gly Asp Gly Leu Phe His Glu Val Ile Asn Gly Leu Gly Glu Arg Asp

65 70 75 80

Asp Tyr Leu Glu Ala Phe Lys Leu Pro Val Cys Met Ile Pro Gly Gly

85 90 95

Ser Gly Asn Ala Phe Ser Tyr Asn Ala Thr Gly Gln Leu Lys Pro Ala

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120

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Met Thr Phe Glu

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<211> 138

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Glu Thr Phe Ala Asn Thr Val Gly Pro Lys Leu Asp Lys Ser Leu Ile

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Arg Tyr Glu Val Val Val Thr Thr Gly Pro Asn His Ala Arg Asn Val

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45

Leu Met Thr Lys Ala Asp Leu Gly Lys Phe Asn Gly Val Leu Ile Leu

50

55

60

Ser Gly Asp Gly Leu Val Phe Glu Ala Leu Asn Gly Ile Leu Cys Arg

65

70

75

80

Glu Asp Ala Phe Arg Ile Phe Pro Thr Leu Pro Ile Gly Ile Val Pro

85

90

95

- 13 -

Ser Gly Ser Gly Asn Gly Leu Leu Cys Ser Val Leu Ser Lys Tyr Gly
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Thr Lys Met Asn Glu Lys Ser Val Met Glu Arg Ala Leu Glu Ile Ala
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Thr Ser Pro Thr Ala Lys Ala Glu Ser Val
130 135

<210> 12

<211> 137

<212> PRT

<213> Arabidopsis

<400> 12

Arg Leu Leu Val Phe Val Asn Pro Phe Gly Gly Lys Lys Ser Ala Arg
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Glu Ile Phe Val Lys Glu Val Lys Pro Leu Phe Glu Asp Ala Asp Val
20 25 30

Gln Leu Glu Ile Gln Glu Thr Lys Tyr Gln Leu His Ala Lys Glu Phe
35 40 45

Val Lys Ser Met Asp Val Ser Lys Tyr Asp Gly Ile Val Cys Val Ser
50 55 60

Gly Asp Gly Ile Leu Val Glu Val Val Asn Gly Leu Leu Glu Arg Ala
65 70 75 80

- 14 -

Asp Trp Arg Asn Ala Leu Lys Leu Pro Ile Gly Met Val Pro Ala Gly
85 90 95

Thr Gly Asn Gly Met Ile Lys Ser Leu Leu Asp Thr Val Gly Leu Arg
100 105 110

Cys Cys Ala Asn Ser Ala Thr Ile Ser Ile Ile Arg Gly His Lys Arg
115 120 125

Ser Val Asp Val Ala Thr Ile Ala Gln
130 135

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tcatgtctga cgacggcctg atgcac

26

<210> 21

<211> 23

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gtctggagat gcattgatgc acg

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gccatccagc ggccgctgtg tagc

24

<210> 25

<211> 23

<212> DNA

<213> mammalian

<400> 25

agcctccctg cagcctcttg caa

23

<210> 26

<211> 19

<212> DNA

<213> mammalian

<400> 26

tcccagcaga ctctggcaa

19

- 18 -

<210> 27

<211> 24

<212> DNA

<213> mammalian

<400> 27

cccagcagga tccgacaacg cgct

24

INTERNATIONAL SEARCH REPORT

International application No.

PCT/AU01/00730

A. CLASSIFICATION OF SUBJECT MATTER		
Int. Cl. ⁷ : C12N 15/54, 9/10; C12Q 1/48; A61K 38/45.		
According to International Patent Classification (IPC) or to both national classification and IPC		
B. FIELDS SEARCHED		
Minimum documentation searched (classification system followed by classification symbols) WORLD PATENT INDEX (WPI) AND CHEMICAL ABSTRACTS (CA): Keywords (KW) see electronic database box below.		
Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched MEDLINE (MD) AND DERWENT BIOTECHNOLOGY ABSTRACTS (DB)): Keywords (KW) see below. :		
Electronic data base consulted during the international search (name of data base and, where practicable, search terms used) KW for WPI and MD: sphingosine kinase; for DB and CA: sphingosine kinase and (mutat? or varian? or ablat? or reduc? or decreas? or lower?).		
C. DOCUMENTS CONSIDERED TO BE RELEVANT		
Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
P,X	WO 01/31029, A (WARNER-LAMBERT COMPANY) published 3 May 2001. See the whole document, especially example 2.	1-7, 9-15 and 17.
P,X	WO 00/70028, A (JOHNSON & JOHNSON RESEARCH PTY LIMITED) published 23 November 2000. See the whole document. especially example 8.	1-7, 9-15 and 17.
X	WO 99/61581, A (OFFICE OF THE DEAN OF RESEARCH AND GRADUATE EDUCATION) published 2 December 1999. See the whole document, especially page 19 and example 1.	1-7, 9-15 and 17.
<input checked="" type="checkbox"/> Further documents are listed in the continuation of Box C <input checked="" type="checkbox"/> See patent family annex		
* Special categories of cited documents:	"T" later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention "X" document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step when the document is taken alone "Y" document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art "&" document member of the same patent family	
"A" document defining the general state of the art which is not considered to be of particular relevance		
"E" earlier application or patent but published on or after the international filing date		
"L" document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified)		
"O" document referring to an oral disclosure, use, exhibition or other means		
"P" document published prior to the international filing date but later than the priority date claimed		
Date of the actual completion of the international search 16 August 2001	Date of mailing of the international search report 21 August 2001	
Name and mailing address of the ISA/AU AUSTRALIAN PATENT OFFICE PO BOX 200, WODEN ACT 2606, AUSTRALIA E-mail address: pct@ipaustalia.gov.au Facsimile No. (02) 6285 3929	Authorized officer J.H. CHAN Telephone No : (02) 6283 2340	

INTERNATIONAL SEARCH REPORT

International application No.

PCT/AU01/00730

C (Continuation). DOCUMENTS CONSIDERED TO BE RELEVANT		
Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
P,X	Pitson S M <i>et al</i> "Expression of a catalytically inactive sphingosine kinase mutant blocks agonist-induced sphingosine kinase activation. A dominant-negative sphingosine kinase". J Biol Chem, 27 October 2000, 275(43) pp33945-33950. See the whole document.	1-32.
P,X	Liu H <i>et al</i> "Molecular cloning and functional characterization of a novel mammalian sphingosine kinase type 2 isoform". J Biol Chem, 30 June 2000, 275(26) pp 19513-19520. See the whole document, especially pages 19515-19516 and figure 1.	1-7, 9-15 and 17.
PX	Pitson S M <i>et al</i> "Human sphingosine kinase: purification, molecular cloning and characterization of the native and recombinant enzymes". Biochem J (2000) 350 part 2:429-441 (published on 1 st September). See the whole document, especially page 436.	1-7, 9-15 and 17.
X	Nava V E <i>et al</i> "Functional characterization of human sphingosine kinase-1". FEBS Lett, 4 May 2000, 473(1):81-84. See the whole document. especially page 82.	1-7, 9-15 and 17.
X	Melendez A J <i>et al</i> "Human sphingosine kinase: molecular cloning, functional characterization and tissue distribution". Gene, 13 June 2000, 251(1):19-26. See the whole document. especially page 23.	1-7, 9-15 and 17.
X	Kohama T <i>et al</i> "Molecular cloning and functional characterization of murine sphingosine kinase". J Biol Chem 273(37) pp23722-23728 (1998) See the whole document. especially pages 23724-23728.	1-7, 9-15 and 17.
A	Topham M K and Prescott S M "Mammalian diacylglycerol kinases, a family of lipid kinases with signaling functions". J Biol Chem 274(17) pp 11447-11450 (1999).	

INTERNATIONAL SEARCH REPORT

International application No.

PCT/AU01/00730

Box I Observations where certain claims were found unsearchable (Continuation of item 2 of first sheet)

This international search report has not been established in respect of certain claims under Article 17(2)(a) for the following reasons:

1. ☐ Claims Nos :
because they relate to subject matter not required to be searched by this Authority, namely:
2. ☒ Claims Nos : 18
because they relate to parts of the international application that do not comply with the prescribed requirements to such an extent that no meaningful international search can be carried out, specifically:
the scope of the term "FOSK" in the claim is unclear and insufficiently defined in the description as such the ambit of the claim is indefinite. Consequently no meaningful international search can be carried out.
3. ☐ Claims Nos :
because they are dependent claims and are not drafted in accordance with the second and third sentences of Rule 6.4(a)

Box II Observations where unity of invention is lacking (Continuation of item 3 of first sheet)

This International Searching Authority found multiple inventions in this international application, as follows:

1. ☐ As all required additional search fees were timely paid by the applicant, this international search report covers all searchable claims
2. ☐ As all searchable claims could be searched without effort justifying an additional fee, this Authority did not invite payment of any additional fee.
3. ☐ As only some of the required additional search fees were timely paid by the applicant, this international search report covers only those claims for which fees were paid, specifically claims Nos.:
4. ☐ No required additional search fees were timely paid by the applicant. Consequently, this international search report is restricted to the invention first mentioned in the claims; it is covered by claims Nos.:

Remark on Protest

- ☐ The additional search fees were accompanied by the applicant's protest.
- ☐ No protest accompanied the payment of additional search fees.



INTERNATIONAL SEARCH REPORT
Information on patent family members

International application No.
PCT/AU01/00730

This Annex lists the known "A" publication level patent family members relating to the patent documents cited in the above-mentioned international search report. The Australian Patent Office is in no way liable for these particulars which are merely given for the purpose of information.

Patent Document Cited in Search Report		Patent Family Member	
WO	200131029	NONE	
WO	200070028	AU	200045239
WO	9961581	AU	40979/99
END OF ANNEX			